

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201623_s_at	0.00228	gb:BC000629.1 /DEF=Homo sapiens, Similar to aspartyl-tRNA synthetase, clone MGC:1562, mRNA, complete cds. /FEA=mRNA /PROD=Similar to aspartyl-tRNA synthetase /DB_XREF=gi:12653688 /UG=Hs.80758 aspartyl-tRNA synthetase /FL=gb:BC000629.1 gb:J05032.1 gb:NM_001349.1	BC000629		NP_001340
201625_s_at	0.005339	insulin induced gene 1	BE300521	Hs.56205	NP_005533
201626_at	0.004341	insulin induced gene 1	BE300521	Hs.56205	NP_005533
201627_s_at	1.26E-04	gb:NM_005542.1 /DEF=Homo sapiens insulin induced gene 1 (INSIG1), mRNA. /FEA=mRNA /GEN=INSIG1 /PROD=insulin induced gene 1 /DB_XREF=gi:5031800 /UG=Hs.56205 insulin induced gene 1 /FL=gb:NM_005542.1	NM_005542		NP_005533
201628_s_at	1.26E-04	gb:NM_006570.1 /DEF=Homo sapiens Ras-related GTP-binding protein (RAGA), mRNA. /FEA=mRNA /GEN=RAGA /PROD=Ras-related GTP-binding protein /DB_XREF=gi:5729998 /UG=Hs.57304 Ras-related GTP-binding protein /FL=gb:U41654.1 gb:NM_006570.1	NM_006570		NP_006561
201631_s_at	0.005852	gb:NM_003897.1 /DEF=Homo sapiens immediate early response 3 (IER3), mRNA. /FEA=mRNA /GEN=IER3 /PROD=immediate early response 3 /DB_XREF=gi:4503328 /UG=Hs.76095 immediate early response 3 /FL=gb:BC000844.1 gb:BC005080.1 gb:AF083421.1 gb:NM_003897.1	NM_003897		NP_434702
201635_s_at	0.029453	fragile X mental retardation, autosomal homolog 1	AI990766	Hs.82712	NP_005078

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201642_at	3.26E-04	gb:NM_005534.1 /DEF=Homo sapiens interferon gamma receptor 2 (interferon gamma transducer 1) (IFNGR2), mRNA. /FEA=mRNA /GEN=IFNGR2 /PROD=interferon gamma receptor 2 (interferon gammatransducer 1) /DB_XREF=gi:5031782 /UG=Hs.177559 interferon gamma receptor 2 (interferon gamma transducer 1) /FL=gb:BC003624.1 gb:U05875.1 gb:U05877.1 gb:NM_005534.1	NM_005534		NP_005525
201643_x_at	3.26E-04	gb:NM_016604.1 /DEF=Homo sapiens putative zinc finger protein (LOC51780), mRNA. /FEA=mRNA /GEN=LOC51780 /PROD=putative zinc finger protein /DB_XREF=gi:7706598 /UG=Hs.24125 putative zinc finger protein /FL=gb:AF251039.1 gb:NM_016604.1	NM_016604		NP_057688
201646_at	0.022926	scavenger receptor class B, member 2	AA885297	Hs.323567	
201648_at	0.00228	Janus kinase 1 (a protein tyrosine kinase)	AL039831	Hs.50651	
201651_s_at	7.58E-04	gb:NM_007229.1 /DEF=Homo sapiens protein kinase C and casein kinase substrate in neurons 2 (PACSIN2), mRNA. /FEA=mRNA /GEN=PACSIN2 /PROD=protein kinase C and casein kinase substrate inneurons 2 /DB_XREF=gi:6005825 /UG=Hs.18842 protein kinase C and casein kinase substrate in neurons 2 /FL=gb:AF128536.1 gb:NM_007229.1	NM_007229		NP_009160

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201652_at	5.03E-04	gb:NM_006837.1 /DEF=Homo sapiens COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 5 (COPS5), mRNA. /FEA=mRNA /GEN=COPS5 /PROD=COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 5 /DB_XREF=gi:5803045 /UG=Hs.198767 COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 5 /FL=gb:BC001187.1 gb:BC001859.1 gb:U65928.1 gb:U70734.1 gb:NM_006837.1	NM_006837		NP_006828
201662_s_at	0.001116	gb:D89053.1 /DEF=Homo sapiens mRNA for Acyl-CoA synthetase 3, complete cds. /FEA=mRNA /PROD=Acyl-CoA synthetase 3 /DB_XREF=gi:4165017 /UG=Hs.268012 fatty-acid-Coenzyme A ligase, long-chain 3 /FL=gb:NM_004457.2 gb:D89053.1 gb:AF116690.1	D89053		NP_004448
201663_s_at	0.04734	gb:NM_005496.1 /DEF=Homo sapiens chromosome-associated polypeptide C (CAP-C), mRNA. /FEA=mRNA /GEN=CAP-C /PROD=chromosome-associated polypeptide C /DB_XREF=gi:4885112 /UG=Hs.50758 SMC4 (structural maintenance of chromosomes 4, yeast)-like 1 /FL=gb:AB019987.1 gb:NM_005496.1 gb:AL136877.1	NM_005496		NP_005487

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201666_at	1.26E-04	gb:NM_003254.1 /DEF=Homo sapiens tissue inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor) (TIMP1), mRNA. /FEA=mRNA /GEN=TIMP1 /PROD=tissue inhibitor of metalloproteinase 1precursor /DB_XREF=gi:4507508 /UG=Hs.5831 tissue inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor) /FL=gb:BC000866.1 gb:M12670.1 gb:M59906.1 gb:NM_003254.1	NM_003254		NP_003245
201669_s_at	1.26E-04	gb:NM_002356.4 /DEF=Homo sapiens myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L) (MACS), mRNA. /FEA=mRNA /GEN=MACS /PROD=myristoylated alanine-rich protein kinase Csubstrate /DB_XREF=gi:11125771 /UG=Hs.75607 myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L) /FL=gb:NM_002356.4 gb:M68956.1 gb:D10522.1	NM_002356		NP_002347
201670_s_at	3.26E-04	gb:M68956.1 /DEF=Human myristoylated alanine-rich C-kinase substrate mRNA, complete cds. /FEA=mRNA /GEN=MACS /PROD=myristoylated alanine-rich C-kinase substrate /DB_XREF=gi:187386 /UG=Hs.75607 myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L) /FL=gb:NM_002356.4 gb:M68956.1 gb:D10522.1	M68956		NP_002347

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201671_x_at	0.011152	gb:BC003556.1 /DEF=Homo sapiens, ubiquitin specific protease 14 (tRNA-guanine transglycosylase), clone MGC:1453, mRNA, complete cds. /FEA=mRNA /PROD=ubiquitin specific protease 14 (tRNA-guaninetransglycosylase) /DB_XREF=gi:13097695 /UG=Hs.75981 ubiquitin specific protease 14 (tRNA-guanine transglycosylase) /FL=gb:BC003556.1 gb:NM_005151.1 gb:U30888.1	BC003556		NP_005142
201673_s_at	0.033381	gb:NM_002103.1 /DEF=Homo sapiens glycogen synthase 1 (muscle) (GYS1), mRNA. /FEA=mRNA /GEN=GYS1 /PROD=glycogen synthase 1 (muscle) /DB_XREF=gi:4504232 /UG=Hs.772 glycogen synthase 1 (muscle) /FL=gb:U32573.1 gb:BC002617.1 gb:J04501.1 gb:NM_002103.1	NM_002103		NP_002094
201684_s_at	5.03E-04	KIAA0737 gene product	BE783632	Hs.194035	NP_055643
201688_s_at	0.007052	tumor protein D52	BE974098	Hs.2384	NP_005070
201690_s_at	0.009689	tumor protein D52	BE974098	Hs.2384	NP_005070
201695_s_at	2.05E-04	gb:NM_000270.1 /DEF=Homo sapiens nucleoside phosphorylase (NP), mRNA. /FEA=mRNA /GEN=NP /PROD=purine nucleoside phosphorylase /DB_XREF=gi:4557800 /UG=Hs.75514 nucleoside phosphorylase /FL=gb:NM_000270.1	NM_000270		NP_000261
201698_s_at	1.26E-04	gb:NM_003769.1 /DEF=Homo sapiens splicing factor, arginineserine-rich 9 (SFRS9), mRNA. /FEA=mRNA /GEN=SFRS9 /PROD=splicing factor, arginineserine-rich 9 /DB_XREF=gi:4506902 /UG=Hs.77608 splicing factor, arginineserine-rich 9 /FL=gb:U30825.1 gb:NM_003769.1	NM_003769		NP_003760

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201699_at	0.003171	gb:NM_002806.1 /DEF=Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 6 (PSMC6), mRNA. /FEA=mRNA /GEN=PSMC6 /PROD=proteasome (prosome, macropain) 26S subunit,ATPase, 6 /DB_XREF=gi:4506214 /UG=Hs.79357 proteasome (prosome, macropain) 26S subunit, ATPase, 6 /FL=gb:BC005390.1 gb:D78275.1 gb:AF006305.1 gb:NM_002806.1	NM_002806		NP_002797
201700_at	0.013223	gb:NM_001760.1 /DEF=Homo sapiens cyclin D3 (CCND3), mRNA. /FEA=mRNA /GEN=CCND3 /PROD=cyclin D3 /DB_XREF=gi:4502618 /UG=Hs.83173 cyclin D3 /FL=gb:M90814.1 gb:M92287.1 gb:NM_001760.1	NM_001760		NP_001751
201703_s_at	6.79E-04	gb:NM_002714.1 /DEF=Homo sapiens protein phosphatase 1, regulatory subunit 10 (PPP1R10), mRNA. /FEA=mRNA /GEN=PPP1R10 /PROD=protein phosphatase 1, regulatory subunit 10 /DB_XREF=gi:4506008 /UG=Hs.106019 protein phosphatase 1, regulatory subunit 10 /FL=gb:NM_002714.1	NM_002714		NP_002705
201705_at	0.001116	gb:NM_002811.1 /DEF=Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 7 (Mov34 homolog) (PSMD7), mRNA. /FEA=mRNA /GEN=PSMD7 /PROD=proteasome (prosome, macropain) 26S subunit,non-ATPase, 7 (Mov34 homolog) /DB_XREF=gi:4506230 /UG=Hs.155543 proteasome (prosome, macropain) 26S subunit, non-ATPase, 7 (Mov34 homolog) /FL=gb:NM_002811.1 gb:D50063.1	NM_002811		NP_002802

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201709_s_at	0.046687	gb:NM_003634.1 /DEF=Homo sapiens NIPSNAP, C. elegans, homolog 1 (NIPSNAP1), mRNA. /FEA=mRNA /GEN=NIPSNAP1 /PROD=NIPSNAP, C. elegans, homolog 1 /DB_XREF=gi:4505398 /UG=Hs.173878 NIPSNAP, C. elegans, homolog 1 /FL=gb:BC002371.1 gb:NM_003634.1	NM_003634		NP_003625
201713_s_at	0.00228	gb:D42063.1 /DEF=Human mRNA for RanBP2 (Ran-binding protein 2), complete cds. /FEA=mRNA /PROD=RanBP2 (Ran-binding protein 2) /DB_XREF=gi:924266 /UG=Hs.199179 RAN binding protein 2 /FL=gb:NM_006267.2 gb:D42063.1	D42063		NP_006258
201716_at	1.26E-04	gb:NM_003099.1 /DEF=Homo sapiens sorting nexin 1 (SNX1), mRNA. /FEA=mRNA /GEN=SNX1 /PROD=sorting nexin 1 /DB_XREF=gi:4507138 /UG=Hs.75283 sorting nexin 1 /FL=gb:BC000357.1 gb:U53225.1 gb:AF065483.1 gb:NM_003099.1	NM_003099		NP_690039
201722_s_at	1.26E-04	Consensus includes gb:AV692127 /FEA=EST /DB_XREF=gi:10293990 /DB_XREF=est:AV692127 /CLONE=GKCAOB04 /UG=Hs.80120 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1) /FL=gb:U41514.1 gb:NM_020474.2	NM_020474		NP_065207

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201724_s_at	3.26E-04	gb:NM_020474.2 /DEF=Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1) (GALNT1), mRNA. /FEA=mRNA /GEN=GALNT1 /PROD=polypeptide N-acetylgalactosaminyltransferase 1 /DB_XREF=gi:13124890 /UG=Hs.80120 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1) /FL=gb:U41514.1 gb:NM_020474.2	NM_020474		NP_065207
201725_at	3.26E-04	gb:NM_006023.1 /DEF=Homo sapiens D123 gene product (D123), mRNA. /FEA=mRNA /GEN=D123 /PROD=D123 gene product /DB_XREF=gi:5174422 /UG=Hs.82043 D123 gene product /FL=gb:BC001600.1 gb:D14878.1 gb:U27112.1 gb:NM_006023.1	NM_006023		NP_006014
201726_at	0.033381	gb:BC003376.1 /DEF=Homo sapiens, ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R), clone MGC:5084, mRNA, complete cds. /FEA=mRNA /PROD=ELAV (embryonic lethal, abnormal vision,Drosophila)-like 1 (Hu antigen R) /DB_XREF=gi:13097227 /UG=Hs.12379 ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R) /FL=gb:U38175.1 gb:BC003376.1 gb:NM_001419.1	BC003376		NP_001410
201730_s_at	0.007779	translocated promoter region (to activated MET oncogene)	BF110993	Hs.169750	NP_003283
201736_s_at	0.02145	similar to S. cerevisiae SSM4	BF000409	Hs.380875	NP_005876

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201737_s_at	0.026891	gb:NM_005885.1 /DEF=Homo sapiens similar to S. cerevisiae SSM4 (TEB4), mRNA. /FEA=mRNA /GEN=TEB4 /PROD=similar to S. cerevisiae SSM4 /DB_XREF=gi:5032166 /UG=Hs.20141 similar to S. cerevisiae SSM4 /FL=gb:AF009301.1 gb:NM_005885.1	NM_005885		NP_005876
201738_at	1.26E-04	gb:NM_005875.1 /DEF=Homo sapiens translation factor sui1 homolog (GC20), mRNA. /FEA=mRNA /GEN=GC20 /PROD=translation factor sui1 homolog /DB_XREF=gi:5031710 /UG=Hs.21756 translation factor sui1 homolog /FL=gb:AF064607.1 gb:AF077052.1 gb:NM_005875.1 gb:AF263452.1	NM_005875		NP_005866
201743_at	0.033381	gb:NM_000591.1 /DEF=Homo sapiens CD14 antigen (CD14), mRNA. /FEA=mRNA /GEN=CD14 /PROD=CD14 antigen precursor /DB_XREF=gi:4557416 /UG=Hs.75627 CD14 antigen /FL=gb:M86511.1 gb:AF097942.1 gb:NM_000591.1	NM_000591		NP_000582
201745_at	0.003171	gb:NM_002822.1 /DEF=Homo sapiens protein tyrosine kinase 9 (PTK9), mRNA. /FEA=mRNA /GEN=PTK9 /PROD=protein tyrosine kinase 9 /DB_XREF=gi:4506274 /UG=Hs.82643 protein tyrosine kinase 9 /FL=gb:NM_002822.1 gb:U02680.1	NM_002822		NP_002813
201758_at	1.26E-04	gb:NM_006292.1 /DEF=Homo sapiens tumor susceptibility gene 101 (TSG101), mRNA. /FEA=mRNA /GEN=TSG101 /PROD=tumor susceptibility gene 101 /DB_XREF=gi:5454139 /UG=Hs.118910 tumor susceptibility gene 101 /FL=gb:BC002487.1 gb:U82130.1 gb:NM_006292.1	NM_006292		NP_006283

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201760_s_at	1.26E-04	gb:NM_018639.1 /DEF=Homo sapiens CS box-containing WD protein (LOC55884), mRNA. /FEA=mRNA /GEN=LOC55884 /PROD=CS box-containing WD protein /DB_XREF=gi:8923880 /UG=Hs.136644 CS box-containing WD protein /FL=gb:AF229181.1 gb:AF163324.1 gb:NM_018639.1	NM_018639		NP_061109
201761_at	7.58E-04	gb:NM_006636.2 /DEF=Homo sapiens methylene tetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase (MTHFD2), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=MTHFD2 /PROD=methylene tetrahydrofolate dehydrogenase (NAD+dependent), methenyltetrahydrofolate cyclohydrolase,precursor /DB_XREF=gi:13699869 /UG=Hs.154672 methylene tetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase /FL=gb:NM_006636.2	NM_006636		NP_006627
201765_s_at	0.023856	hexosaminidase A (alpha polypeptide)	AL523158	Hs.119403	NP_000511
201769_at	1.26E-04	gb:NM_014666.1 /DEF=Homo sapiens KIAA0171 gene product (KIAA0171), mRNA. /FEA=mRNA /GEN=KIAA0171 /PROD=KIAA0171 gene product /DB_XREF=gi:7661967 /UG=Hs.155623 KIAA0171 gene product /FL=gb:D79993.1 gb:BC004467.1 gb:NM_014666.1	NM_014666		NP_055481

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201770_at	0.045608	gb:NM_004596.1 /DEF=Homo sapiens small nuclear ribonucleoprotein polypeptide A (SNRPA), mRNA. /FEA=mRNA /GEN=SNRPA /PROD=small nuclear ribonucleoprotein polypeptide A /DB_XREF=gi:4759155 /UG=Hs.173255 small nuclear ribonucleoprotein polypeptide A /FL=gb:BC000405.1 gb:NM_004596.1	NM_004596		NP_004587
201772_at	7.58E-04	gb:NM_015878.1 /DEF=Homo sapiens antizyme inhibitor (LOC51582), mRNA. /FEA=mRNA /GEN=LOC51582 /PROD=antizyme inhibitor /DB_XREF=gi:7706219 /UG=Hs.223014 antizyme inhibitor /FL=gb:D88674.1 gb:NM_015878.1	NM_015878		NP_680479
201773_at	7.58E-04	gb:NM_015339.1 /DEF=Homo sapiens activity-dependent neuroprotective protein (ADNP), mRNA. /FEA=mRNA /GEN=ADNP /PROD=activity-dependent neuroprotective protein /DB_XREF=gi:12229216 /UG=Hs.3657 activity-dependent neuroprotective protein /FL=gb:AF250860.1 gb:NM_015339.1	NM_015339		NP_056154
201776_s_at	0.005852	Consensus includes gb:AK001487.1 /DEF=Homo sapiens cDNA FLJ10625 fis, clone NT2RP2005540, highly similar to Homo sapiens mRNA for KIAA0494 protein. /FEA=mRNA /DB_XREF=gi:7022773 /UG=Hs.62515 KIAA0494 gene product /FL=gb:BC002525.1 gb:AB007963.1 gb:NM_014774.1	AK001487		NP_055589

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201780_s_at	1.26E-04	gb:NM_007282.1 /DEF=Homo sapiens ring finger protein 13 (RNF13), mRNA. /FEA=mRNA /GEN=RNF13 /PROD=ring finger protein 13 /DB_XREF=gi:6005863 /UG=Hs.6900 ring finger protein 13 /FL=gb:AF037204.1 gb:AF070558.1 gb:NM_007282.1	NM_007282		NP_009213
201784_s_at	0.003171	gb:NM_014267.1 /DEF=Homo sapiens small acidic protein (IMAGE145052), mRNA. /FEA=mRNA /GEN=IMAGE145052 /PROD=small acidic protein /DB_XREF=gi:7657233 /UG=Hs.78050 small acidic protein /FL=gb:U51678.1 gb:NM_014267.1	NM_014267		NP_055082
201794_s_at	0.010205	gb:NM_014837.1 /DEF=Homo sapiens KIAA0250 gene product (KIAA0250), mRNA. /FEA=mRNA /GEN=KIAA0250 /PROD=KIAA0250 gene product /DB_XREF=gi:7662023 /UG=Hs.15087 KIAA0250 gene product /FL=gb:D87437.1 gb:NM_014837.1	NM_014837		NP_055652
201795_at	0.004341	gb:NM_002296.1 /DEF=Homo sapiens lamin B receptor (LBR), mRNA. /FEA=mRNA /GEN=LBR /PROD=lamin B receptor /DB_XREF=gi:4504960 /UG=Hs.152931 lamin B receptor /FL=gb:L25931.1 gb:NM_002296.1	NM_002296		NP_002287
201798_s_at	0.026891	gb:NM_013451.1 /DEF=Homo sapiens fer-1 (C.elegans)-like 3 (myoferlin) (FER1L3), mRNA. /FEA=mRNA /GEN=FER1L3 /PROD=fer-1 (C.elegans)-like 3 (myoferlin) /DB_XREF=gi:7305052 /UG=Hs.234680 fer-1 (C.elegans)-like 3 (myoferlin) /FL=gb:AF182316.1 gb:NM_013451.1	NM_013451		NP_579899

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201805_at	0.003171	gb:NM_002733.1 /DEF=Homo sapiens protein kinase, AMP-activated, gamma 1 non-catalytic subunit (PRKAG1), mRNA. /FEA=mRNA /GEN=PRKAG1 /PROD=protein kinase, AMP-activated, gamma 1non-catalytic subunit /DB_XREF=gi:4506060 /UG=Hs.3136 protein kinase, AMP-activated, gamma 1 non-catalytic subunit /FL=gb:BC000358.1 gb:U42412.1 gb:NM_002733.1	NM_002733		NP_002724
201807_at	3.26E-04	gb:NM_004896.1 /DEF=Homo sapiens vacuolar protein sorting 26 (yeast homolog) (VPS26), mRNA. /FEA=mRNA /GEN=VPS26 /PROD=vacuolar protein sorting 26 (yeast homolog) /DB_XREF=gi:4758509 /UG=Hs.67052 vacuolar protein sorting 26 (yeast homolog) /FL=gb:AF054179.1 gb:NM_004896.1 gb:AF175266.1	NM_004896		NP_004887
201810_s_at	0.033381	SH3-domain binding protein 5 (BTK-associated)	AL562152	Hs.109150	NP_004835
201811_x_at	2.05E-04	gb:NM_004844.1 /DEF=Homo sapiens SH3-domain binding protein 5 (BTK-associated) (SH3BP5), mRNA. /FEA=mRNA /GEN=SH3BP5 /PROD=SH3-domain binding protein 5 (BTK-associated) /DB_XREF=gi:4759057 /UG=Hs.109150 SH3-domain binding protein 5 (BTK-associated) /FL=gb:AB005047.1 gb:NM_004844.1	NM_004844		NP_004835
201814_at	0.007779	Consensus includes gb:AI300084 /FEA=EST /DB_XREF=gi:3959430 /DB_XREF=est:qn59f02.x1 /CLONE=IMAGE:1902555 /UG=Hs.115740 KIAA0210 gene product /FL=gb:D86965.1 gb:NM_014744.1	NM_014744		NP_055559

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201815_s_at	0.028685	gb:NM_014744.1 /DEF=Homo sapiens KIAA0210 gene product (KIAA0210), mRNA. /FEA=mRNA /GEN=KIAA0210 /PROD=KIAA0210 gene product /DB_XREF=gi:7661997 /UG=Hs.115740 KIAA0210 gene product /FL=gb:D86965.1 gb:NM_014744.1	NM_014744		NP_055559
201816_s_at	0.001116	gb:NM_001483.1 /DEF=Homo sapiens glioblastoma amplified sequence (GBAS), mRNA. /FEA=mRNA /GEN=GBAS /PROD=glioblastoma amplified sequence /DB_XREF=gi:4503936 /UG=Hs.152707 glioblastoma amplified sequence /FL=gb:BC000732.1 gb:BC001837.1 gb:AF029786.1 gb:NM_001483.1	NM_001483		NP_001474
201817_at	3.26E-04	gb:NM_014671.1 /DEF=Homo sapiens KIAA0010 gene product (KIAA0010), mRNA. /FEA=mRNA /GEN=KIAA0010 /PROD=KIAA0010 gene product /DB_XREF=gi:7661855 /UG=Hs.155287 KIAA0010 gene product /FL=gb:D13635.1 gb:NM_014671.1	NM_014671		NP_055486
201818_at	0.033381	gb:NM_024830.1 /DEF=Homo sapiens hypothetical protein FLJ12443 (FLJ12443), mRNA. /FEA=mRNA /GEN=FLJ12443 /PROD=hypothetical protein FLJ12443 /DB_XREF=gi:13376233 /UG=Hs.179882 hypothetical protein FLJ12443 /FL=gb:NM_024830.1	NM_024830		NP_079106

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201821_s_at	0.009716	gb:BC004439.1 /DEF=Homo sapiens, translocase of inner mitochondrial membrane 17 (yeast) homolog A, clone MGC:4031, mRNA, complete cds. /FEA=mRNA /PROD=translocase of inner mitochondrial membrane 17(yeast) homolog A /DB_XREF=gi:13325239 /UG=Hs.20716 translocase of inner mitochondrial membrane 17 (yeast) homolog A /FL=gb:BC004439.1 gb:AF106622.1 gb:NM_006335.1	BC004439		NP_006326
201823_s_at	0.001116	gb:NM_004290.1 /DEF=Homo sapiens ring finger protein 14 (RNF14), mRNA. /FEA=mRNA /GEN=RNF14 /PROD=ring finger protein 14 /DB_XREF=gi:4757761 /UG=Hs.215857 ring finger protein 14 /FL=gb:AF060544.1 gb:NM_004290.1 gb:AB022663.1	NM_004290		NP_004281
201824_at	0.010205	gb:AB022663.1 /DEF=Homo sapiens HFB30 mRNA, complete cds. /FEA=mRNA /GEN=HFB30 /DB_XREF=gi:5019617 /UG=Hs.215857 ring finger protein 14 /FL=gb:AF060544.1 gb:NM_004290.1 gb:AB022663.1	AB022663		NP_004281
201826_s_at	0.004733	gb:NM_016002.1 /DEF=Homo sapiens CGI-49 protein (LOC51097), mRNA. /FEA=mRNA /GEN=LOC51097 /PROD=CGI-49 protein /DB_XREF=gi:7705766 /UG=Hs.238126 CGI-49 protein /FL=gb:AF151807.1 gb:NM_016002.1	NM_016002		NP_057086
201833_at	0.026891	gb:NM_001527.1 /DEF=Homo sapiens histone deacetylase 2 (HDAC2), mRNA. /FEA=mRNA /GEN=HDAC2 /PROD=histone deacetylase 2 /DB_XREF=gi:4557640 /UG=Hs.3352 histone deacetylase 2 /FL=gb:U31814.1 gb:NM_001527.1	NM_001527		NP_001518

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201840_at	1.26E-04	gb:NM_006156.1 /DEF=Homo sapiens neural precursor cell expressed, developmentally down-regulated 8 (NEDD8), mRNA. /FEA=mRNA /GEN=NEDD8 /PROD=neural precursor cell expressed, developmentally down-regulated 8 /DB_XREF=gi:5453759 /UG=Hs.75512 neural precursor cell expressed, developmentally down-regulated 8 /FL=gb:D23662.1 gb:NM_006156.1	NM_006156		NP_006147
201845_s_at	1.26E-04	gb:AB029551.1 /DEF=Homo sapiens YEAF1 mRNA for YY1 and E4TF1 associated factor 1, complete cds. /FEA=mRNA /GEN=YEAF1 /PROD=YY1 and E4TF1 associated factor 1 /DB_XREF=gi:6714542 /UG=Hs.7910 RING1 and YY1 binding protein /FL=gb:AF179286.1 gb:AB029551.1 gb:NM_012234.1	AB029551		NP_036366
201846_s_at	0.003171	gb:NM_012234.1 /DEF=Homo sapiens RING1 and YY1 binding protein (RYBP), mRNA. /FEA=mRNA /GEN=RYBP /PROD=RING1 and YY1 binding protein /DB_XREF=gi:6912639 /UG=Hs.7910 RING1 and YY1 binding protein /FL=gb:AF179286.1 gb:AB029551.1 gb:NM_012234.1	NM_012234		NP_036366
201847_at	0.02145	gb:NM_000235.1 /DEF=Homo sapiens lipase A, lysosomal acid, cholesterol esterase (Wolman disease) (LIPA), mRNA. /FEA=mRNA /GEN=LIPA /PROD=lipase A precursor /DB_XREF=gi:4557720 /UG=Hs.85226 lipase A, lysosomal acid, cholesterol esterase (Wolman disease) /FL=gb:M74775.1 gb:NM_000235.1 gb:U08464.1	NM_000235		NP_000226

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201848_s_at	0.03002	gb:U15174.1 /DEF=Homo sapiens BCL2adenovirus E1B 19kD-interacting protein 3 (BNIP3) mRNA, complete cds. /FEA=mRNA /GEN=BNIP3 /PROD=BCL2adenovirus E1B 19kD-interacting protein 3 /DB_XREF=gi:558845 /UG=Hs.79428 BCL2adenovirus E1B 19kD-interacting protein 3 /FL=gb:AF002697.1 gb:U15174.1 gb:NM_004052.2	U15174		NP_004043
201850_at	0.007779	gb:NM_001747.1 /DEF=Homo sapiens capping protein (actin filament), gelsolin-like (CAPG), mRNA. /FEA=mRNA /GEN=CAPG /PROD=capping protein (actin filament), gelsolin-like /DB_XREF=gi:4502560 /UG=Hs.82422 capping protein (actin filament), gelsolin-like /FL=gb:BC000728.1 gb:M94345.1 gb:NM_001747.1	NM_001747		NP_001738
201853_s_at	0.033381	gb:NM_021873.1 /DEF=Homo sapiens cell division cycle 25B (CDC25B), transcript variant 3, mRNA. /FEA=mRNA /GEN=CDC25B /PROD=cell division cycle 25B, isoform 3 /DB_XREF=gi:11641412 /UG=Hs.153752 cell division cycle 25B /FL=gb:NM_021873.1	NM_021873		NP_068660
201857_at	0.013223	gb:NM_016107.1 /DEF=Homo sapiens M-phase phosphoprotein homolog (LOC51663), mRNA. /FEA=mRNA /GEN=LOC51663 /PROD=M-phase phosphoprotein homolog /DB_XREF=gi:7706372 /UG=Hs.173518 M-phase phosphoprotein homolog /FL=gb:BC000376.1 gb:BC000746.1 gb:AF100742.1 gb:NM_016107.1	NM_016107		NP_057191

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201862_s_at	7.58E-04	gb:NM_004735.1 /DEF=Homo sapiens leucine rich repeat (in FLII) interacting protein 1 (LRRFIP1), mRNA. /FEA=mRNA /GEN=LRRFIP1 /PROD=leucine rich repeat (in FLII) interactingprotein 1 /DB_XREF=gi:4758689 /UG=Hs.326159 leucine rich repeat (in FLII) interacting protein 1 /FL=gb:NM_004735.1	NM_004735		NP_004726
201863_at	7.58E-04	gb:NM_014077.1 /DEF=Homo sapiens DKFZP586O0120 protein (DKFZP586O0120), mRNA. /FEA=mRNA /GEN=DKFZP586O0120 /PROD=DKFZP586O0120 protein /DB_XREF=gi:7661695 /UG=Hs.4766 DKFZP586O0120 protein /FL=gb:BC000639.1 gb:AL050157.1 gb:AF151902.1 gb:NM_014077.1	NM_014077		NP_054796
201865_x_at	0.00161	nuclear receptor subfamily 3, group C, member 1 (glucocorticoid receptor)	AI432196	Hs.75772	NP_000167
201866_s_at	7.58E-04	gb:NM_000176.1 /DEF=Homo sapiens nuclear receptor subfamily 3, group C, member 1 (NR3C1), mRNA. /FEA=mRNA /GEN=NR3C1 /PROD=nuclear receptor subfamily 3, group C, member 1 /DB_XREF=gi:4504132 /UG=Hs.75772 nuclear receptor subfamily 3, group C, member 1 /FL=gb:M10901.1 gb:NM_000176.1	NM_000176		NP_000167
201867_s_at	0.008036	Consensus includes gb:AW968555 /FEA=EST /DB_XREF=gi:8158396 /DB_XREF=est:EST380631 /UG=Hs.76536 transducin (beta)-like 1 /FL=gb:NM_005647.1	NM_005647		NP_005638

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201870_at	0.016074	gb:NM_006809.1 /DEF=Homo sapiens translocase of outer mitochondrial membrane 34 (TOM34), mRNA. /FEA=mRNA /GEN=TOM34 /PROD=translocase of outer mitochondrial membrane 34 /DB_XREF=gi:5803204 /UG=Hs.76927 translocase of outer mitochondrial membrane 34 /FL=gb:BC001763.1 gb:U58970.1 gb:NM_006809.1	NM_006809		NP_006800
201879_at	0.026891	Consensus includes gb:AI694332 /FEA=EST /DB_XREF=gi:4971672 /DB_XREF=est:wd45e11.x1 /CLONE=IMAGE:2331116 /UG=Hs.181461 ariadne (Drosophila) homolog, ubiquitin-conjugating enzyme E2-binding protein, 1 /FL=gb:AF072832.1 gb:NM_005744.2	NM_005744		NP_005735
201883_s_at	0.021448	gb:D29805.1 /DEF=Human mRNA for beta-1,4-galactosyltransferase, complete cds. /FEA=mRNA /PROD=beta-1,4-galactosyltransferase /DB_XREF=gi:474986 /UG=Hs.198248 UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 1 /FL=gb:NM_001497.1 gb:D29805.1	D29805		NP_001488
201886_at	0.007052	gb:NM_025230.1 /DEF=Homo sapiens hypthetical protein PRO2389 (PRO2389), mRNA. /FEA=mRNA /GEN=PRO2389 /PROD=hypthetical protein PRO2389 /DB_XREF=gi:13489107 /UG=Hs.283976 hypthetical protein PRO2389 /FL=gb:AF130070.1 gb:AF267858.1 gb:NM_025230.1	NM_025230		NP_079506

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201890_at	0.008237	Consensus includes gb:BE966236 /FEA=EST /DB_XREF=gi:11771437 /DB_XREF=est:601660172R1 /CLONE=IMAGE:3905920 /UG=Hs.75319 ribonucleotide reductase M2 polypeptide /FL=gb:NM_001034.1	NM_001034		NP_001025
201892_s_at	0.002418	gb:NM_000884.1 /DEF=Homo sapiens IMP (inosine monophosphate) dehydrogenase 2 (IMPDH2), mRNA. /FEA=mRNA /GEN=IMPDH2 /PROD=IMP (inosine monophosphate) dehydrogenase 2 /DB_XREF=gi:4504688 /UG=Hs.75432 IMP (inosine monophosphate) dehydrogenase 2 /FL=gb:J04208.1 gb:NM_000884.1	NM_000884		NP_000875
201894_s_at	1.26E-04	gb:NM_001920.1 /DEF=Homo sapiens decorin (DCN), mRNA. /FEA=mRNA /GEN=DCN /PROD=decorin /DB_XREF=gi:4503270 /UG=Hs.76152 decorin /FL=gb:M14219.1 gb:NM_001920.1 gb:AF138300.1 gb:AF138304.1	NM_001920		NP_598014
201895_at	0.003529	gb:NM_001654.1 /DEF=Homo sapiens v-raf murine sarcoma 3611 viral oncogene homolog 1 (ARAF1), mRNA. /FEA=mRNA /GEN=ARAF1 /PROD=v-raf murine sarcoma 3611 viral oncogene homolog1 /DB_XREF=gi:4502192 /UG=Hs.77183 v-raf murine sarcoma 3611 viral oncogene homolog 1 /FL=gb:BC002466.1 gb:NM_001654.1	NM_001654		NP_001645

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201897_s_at	9.08E-04	gb:NM_001826.1 /DEF=Homo sapiens CDC28 protein kinase 1 (CKS1), mRNA. /FEA=mRNA /GEN=CKS1 /PROD=CDC28 protein kinase 1 /DB_XREF=gi:4502856 /UG=Hs.77550 CDC28 protein kinase 1 /FL=gb:BC001425.1 gb:AF274941.1 gb:AF279897.1 gb:NM_001826.1	NM_001826		NP_001817
201898_s_at	1.26E-04	ubiquitin-conjugating enzyme E2A (RAD6 homolog)	AI126625	Hs.80612	NP_003327
201899_s_at	0.00161	gb:NM_003336.1 /DEF=Homo sapiens ubiquitin-conjugating enzyme E2A (RAD6 homolog) (UBE2A), mRNA. /FEA=mRNA /GEN=UBE2A /PROD=ubiquitin-conjugating enzyme E2A (RAD6 homolog) /DB_XREF=gi:4507768 /UG=Hs.80612 ubiquitin-conjugating enzyme E2A (RAD6 homolog) /FL=gb:M74524.1 gb:NM_003336.1	NM_003336		NP_003327
201901_s_at	0.003171	Consensus includes gb:Z14077.1 /DEF=H.sapiens mRNA for YY1NF-E1 protein. /FEA=mRNA /PROD=YY1 NF-E1 /DB_XREF=gi:38010 /UG=Hs.97496 YY1 transcription factor /FL=gb:M77698.1 gb:M76541.1 gb:NM_003403.2	NM_003403		NP_003394
201903_at	3.26E-04	gb:NM_003365.1 /DEF=Homo sapiens ubiquinol-cytochrome c reductase core protein I (UQCRC1), mRNA. /FEA=mRNA /GEN=UQCRC1 /PROD=ubiquinol-cytochrome c reductase core protein I /DB_XREF=gi:4507840 /UG=Hs.119251 ubiquinol-cytochrome c reductase core protein I /FL=gb:L16842.1 gb:NM_003365.1 gb:D26485.1	NM_003365		NP_003356

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201908_at	0.041055	gb:NM_004423.2 /DEF=Homo sapiens dishevelled 3 (homologous to Drosophila dsh) (DVL3), mRNA. /FEA=mRNA /GEN=DVL3 /PROD=dishevelled 3 /DB_XREF=gi:6806886 /UG=Hs.174044 dishevelled 3 (homologous to Drosophila dsh) /FL=gb:U49262.1 gb:D86963.1 gb:U75651.1 gb:AF006013.1 gb:NM_004423.2	NM_004423		NP_004414
201912_s_at	1.26E-04	gb:NM_002094.1 /DEF=Homo sapiens G1 to S phase transition 1 (GSPT1), mRNA. /FEA=mRNA /GEN=GSPT1 /PROD=G1 to S phase transition 1 /DB_XREF=gi:4504166 /UG=Hs.2707 G1 to S phase transition 1 /FL=gb:NM_002094.1	NM_002094		NP_002085
201913_s_at	0.013223	gb:NM_025233.1 /DEF=Homo sapiens nucleotide binding protein (NBP), mRNA. /FEA=mRNA /GEN=NBP /PROD=nucleotide binding protein /DB_XREF=gi:13376837 /UG=Hs.296422 nucleotide binding protein /FL=gb:NM_025233.1 gb:AF208536.1	NM_025233		NP_079509
201914_s_at	0.004341	Consensus includes gb:AK001465.1 /DEF=Homo sapiens cDNA FLJ10603 fis, clone NT2RP2005012, highly similar to Homo sapiens mRNA for SEC63 protein. /FEA=mRNA /DB_XREF=gi:7022740 /UG=Hs.31575 SEC63, endoplasmic reticulum translocon component (S. cerevisiae) like /FL=gb:AF100141.1 gb:NM_007214.1	AK001465		NP_009145

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201915_at	0.007258	Consensus includes gb:AI806665 /FEA=EST /DB_XREF=gi:5393231 /DB_XREF=est:wf35c05.x1 /CLONE=IMAGE:2357576 /UG=Hs.31575 SEC63, endoplasmic reticulum translocon component (S. cerevisiae) like /FL=gb:AF100141.1 gb:NM_007214.1	AK001465		NP_009145
201916_s_at	5.03E-04	gb:NM_007214.1 /DEF=Homo sapiens SEC63, endoplasmic reticulum translocon component (S. cerevisiae) like (SEC63L), mRNA. /FEA=mRNA /GEN=SEC63L /PROD=SEC63, endoplasmic reticulum translocon component (S. cerevisiae) like /DB_XREF=gi:6005871 /UG=Hs.31575 SEC63, endoplasmic reticulum translocon component (S. cerevisiae) like /FL=gb:AF100141.1 gb:NM_007214.1	NM_007214		NP_009145
201917_s_at	1.26E-04	hypothetical protein FLJ10618	AI694452	Hs.408258	NP_060625
201918_at	0.042466	hypothetical protein FLJ10618	AI694452	Hs.408258	NP_060625
201920_at	0.013223	gb:NM_005415.2 /DEF=Homo sapiens solute carrier family 20 (phosphate transporter), member 1 (SLC20A1), mRNA. /FEA=mRNA /GEN=SLC20A1 /PROD=solute carrier family 20 (phosphate transporter), member 1 /DB_XREF=gi:7382462 /UG=Hs.78452 solute carrier family 20 (phosphate transporter), member 1 /FL=gb:L20859.1 gb:NM_005415.2	NM_005415		NP_005406

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201921_at	0.026891	gb:NM_004125.1 /DEF=Homo sapiens guanine nucleotide binding protein 10 (GNG10), mRNA. /FEA=mRNA /GEN=GNG10 /PROD=guanine nucleotide binding protein 10 /DB_XREF=gi:4758445 /UG=Hs.79126 guanine nucleotide binding protein 10 /FL=gb:NM_004125.1 gb:U31383.1	NM_004125		NP_004116
201925_s_at	1.26E-04	gb:NM_000574.1 /DEF=Homo sapiens decay accelerating factor for complement (CD55, Cromer blood group system) (DAF), mRNA. /FEA=mRNA /GEN=DAF /PROD=decay accelerating factor for complement (CD55,Cromer blood group system) /DB_XREF=gi:10835142 /UG=Hs.1369 decay accelerating factor for complement (CD55, Cromer blood group system) /FL=gb:NM_000574.1 gb:BC001288.1 gb:M31516.1	NM_000574		NP_000565
201926_s_at	1.26E-04	gb:BC001288.1 /DEF=Homo sapiens, Similar to decay accelerating factor for complement (CD55, Cromer blood group system), clone MGC:5192, mRNA, complete cds. /FEA=mRNA /PROD=Similar to decay accelerating factor for complement (CD55, Cromer blood group system) /DB_XREF=gi:12654888 /UG=Hs.1369 decay accelerating factor for complement (CD55, Cromer blood group system) /FL=gb:NM_000574.1 gb:BC001288.1 gb:M31516.1	BC001288		NP_000565

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201928_at	0.016934	Consensus includes gb:AA194254 /FEA=EST /DB_XREF=gi:1783969 /DB_XREF=est:zr39d09.s1 /CLONE=IMAGE:665777 /UG=Hs.152151 plakophilin 4 /FL=gb:NM_003628.2	NM_003628		NP_003619
201929_s_at	0.003257	gb:NM_003628.2 /DEF=Homo sapiens plakophilin 4 (PKP4), mRNA. /FEA=mRNA /GEN=PKP4 /PROD=plakophilin 4 /DB_XREF=gi:6806894 /UG=Hs.152151 plakophilin 4 /FL=gb:NM_003628.2	NM_003628		NP_003619
201930_at	3.56E-04	gb:NM_005915.2 /DEF=Homo sapiens minichromosome maintenance deficient (mis5, S. pombe) 6 (MCM6), mRNA. /FEA=mRNA /GEN=MCM6 /PROD=minichromosome maintenance deficient (mis5, S.pombe) 6 /DB_XREF=gi:7427518 /UG=Hs.155462 minichromosome maintenance deficient (mis5, S. pombe) 6 /FL=gb:U46838.1 gb:D84557.1 gb:NM_005915.2	NM_005915		NP_005906
201936_s_at	0.001012	gb:NM_003760.2 /DEF=Homo sapiens eukaryotic translation initiation factor 4 gamma, 3 (EIF4G3), mRNA. /FEA=mRNA /GEN=EIF4G3 /PROD=eukaryotic translation initiation factor 4gamma, 3 /DB_XREF=gi:10092600 /UG=Hs.25732 eukaryotic translation initiation factor 4 gamma, 3 /FL=gb:NM_003760.2 gb:AF012072.2	NM_003760		NP_003751
201940_at	5.03E-04	carboxypeptidase D	AA897514	Hs.5057	NP_001295
201941_at	5.03E-04	carboxypeptidase D	AA897514	Hs.5057	NP_001295

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201943_s_at	0.041055	gb:NM_001304.2 /DEF=Homo sapiens carboxypeptidase D (CPD), mRNA. /FEA=mRNA /GEN=CPD /PROD=carboxypeptidase D precursor /DB_XREF=gi:8051580 /UG=Hs.5057 carboxypeptidase D /FL=gb:U65090.1 gb:D85390.1 gb:NM_001304.2	NM_001304		NP_001295
201944_at	1.26E-04	gb:NM_000521.2 /DEF=Homo sapiens hexosaminidase B (beta polypeptide) (HEXB), mRNA. /FEA=mRNA /GEN=HEXB /PROD=hexosaminidase B preproprotein /DB_XREF=gi:13128866 /UG=Hs.51043 hexosaminidase B (beta polypeptide) /FL=gb:NM_000521.2 gb:M19735.1	NM_000521		NP_000512
201949_x_at	0.007779	capping protein (actin filament) muscle Z-line, beta	AL572341	Hs.333417	NP_004921
201951_at	0.006698	Consensus includes gb:BF242905 /FEA=EST /DB_XREF=gi:11156833 /DB_XREF=est:601877949F1 /CLONE=IMAGE:4106028 /UG=Hs.10247 activated leucocyte cell adhesion molecule /FL=gb:NM_001627.1 gb:L38608.1	NM_001627		NP_001618
201952_at	0.026891	Consensus includes gb:AA156721 /FEA=EST /DB_XREF=gi:1728335 /DB_XREF=est:zl18b04.s1 /CLONE=IMAGE:502255 /UG=Hs.10247 activated leucocyte cell adhesion molecule /FL=gb:NM_001627.1 gb:L38608.1	NM_001627		NP_001618

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201954_at	0.001116	gb:NM_005720.1 /DEF=Homo sapiens actin related protein 23 complex, subunit 1A (41 kD) (ARPC1B), mRNA. /FEA=mRNA /GEN=ARPC1B /PROD=actin related protein 23 complex, subunit 1A(41 kD) /DB_XREF=gi:5031600 /UG=Hs.11538 actin related protein 23 complex, subunit 1A (41 kD) /FL=gb:BC002562.1 gb:AF006084.1 gb:NM_005720.1	NM_005720		NP_005711
201959_s_at	0.001618	KIAA0916 protein	AA488899	Hs.151411	NP_055872
201963_at	0.007779	gb:NM_021122.2 /DEF=Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 2 (FACL2), mRNA. /FEA=mRNA /GEN=FACL2 /PROD=long-chain fatty-acid-Coenzyme A ligase 2 /DB_XREF=gi:12669906 /UG=Hs.154890 fatty-acid-Coenzyme A ligase, long-chain 2 /FL=gb:NM_021122.2 gb:D10040.1	NM_021122		NP_066945
201968_s_at	1.26E-04	gb:NM_002633.1 /DEF=Homo sapiens phosphoglucomutase 1 (PGM1), mRNA. /FEA=mRNA /GEN=PGM1 /PROD=phosphoglucomutase 1 /DB_XREF=gi:4505764 /UG=Hs.1869 phosphoglucomutase 1 /FL=gb:BC001756.1 gb:M83088.1 gb:NM_002633.1	NM_002633		NP_002624
201972_at	0.005852	gb:AF113129.1 /DEF=Homo sapiens vacuolar ATPase isoform VA68 mRNA, complete cds. /FEA=mRNA /PROD=vacuolar ATPase isoform VA68 /DB_XREF=gi:6523820 /UG=Hs.281866 ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump), alpha polypeptide, 70kD, isoform 1 /FL=gb:L09235.1 gb:NM_001690.1 gb:AF113129.1	AF113129		NP_001681
201973_s_at	0.003171	CGI-43 protein	AL550875	Hs.289112	NP_056437

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201975_at	5.03E-04	gb:NM_002956.1 /DEF=Homo sapiens restin (Reed-Steinberg cell-expressed intermediate filament-associated protein) (RSN), mRNA. /FEA=mRNA /GEN=RSN /PROD=restin (Reed-Steinberg cell-expressed intermediate filament-associated protein) /DB_XREF=gi:4506750 /UG=Hs.31638 restin (Reed-Steinberg cell-expressed intermediate filament-associated protein) /FL=gb:NM_002956.1	NM_002956		NP_002947
201980_s_at	0.00228	gb:NM_012425.2 /DEF=Homo sapiens Ras suppressor protein 1 (RSU1), mRNA. /FEA=mRNA /GEN=RSU1 /PROD=ras suppressor protein 1 /DB_XREF=gi:10800408 /UG=Hs.75551 Homo sapiens Ras suppressor protein 1 (RSU1), mRNA /FL=gb:NM_012425.2	NM_012425		NP_036557
201985_at	1.26E-04	gb:NM_014846.1 /DEF=Homo sapiens KIAA0196 gene product (KIAA0196), mRNA. /FEA=mRNA /GEN=KIAA0196 /PROD=KIAA0196 gene product /DB_XREF=gi:7661987 /UG=Hs.8294 KIAA0196 gene product /FL=gb:D83780.1 gb:NM_014846.1	NM_014846		NP_055661
201991_s_at	0.033381	kinesin family member 5B	BF223224	Hs.149436	NP_004512
201995_at	1.26E-04	gb:NM_000127.1 /DEF=Homo sapiens exostoses (multiple) 1 (EXT1), mRNA. /FEA=mRNA /GEN=EXT1 /PROD=exostoses (multiple) 1 /DB_XREF=gi:4557570 /UG=Hs.184161 exostoses (multiple) 1 /FL=gb:BC001174.1 gb:NM_000127.1	NM_000127		NP_000118

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
201997_s_at	0.007779	gb:NM_015001.1 /DEF=Homo sapiens KIAA0929 protein Msx2 interacting nuclear target (MINT) homolog (KIAA0929), mRNA. /FEA=mRNA /GEN=KIAA0929 /PROD=KIAA0929 protein Msx2 interacting nuclear target(MINT) homolog /DB_XREF=gi:7657266 /UG=Hs.184245 KIAA0929 protein Msx2 interacting nuclear target (MINT) homolog /FL=gb:NM_015001.1	NM_015001		NP_055816
201999_s_at	0.007779	gb:NM_006519.1 /DEF=Homo sapiens t-complex-associated-testis-expressed 1-like 1 (TCTEL1), mRNA. /FEA=mRNA /GEN=TCTEL1 /PROD=t-complex-associated-testis-expressed 1-like 1 /DB_XREF=gi:5730084 /UG=Hs.266940 t-complex-associated-testis-expressed 1-like 1 /FL=gb:U56255.1 gb:D50663.1 gb:NM_006519.1	NM_006519		NP_006510
202000_at	0.007779	gb:BC002772.1 /DEF=Homo sapiens, NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14), clone MGC:3686, mRNA, complete cds. /FEA=mRNA /PROD=NADH dehydrogenase (ubiquinone) 1 alphasubcomplex, 6 (14kD, B14) /DB_XREF=gi:12803858 /UG=Hs.274416 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14) /FL=gb:BC002772.1 gb:AF047182.1 gb:NM_002490.1	BC002772		NP_002481

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
202001_s_at	0.026891	gb:NM_002490.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14) (NDUFA6), mRNA. /FEA=mRNA /GEN=NDUFA6 /PROD=NADH dehydrogenase (ubiquinone) 1 alphasubcomplex, 6 (14kD, B14) /DB_XREF=gi:4505358 /UG=Hs.274416 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14) /FL=gb:BC002772.1 gb:AF047182.1 gb:NM_002490.1	NM_002490		NP_002481
202004_x_at	0.016934	gb:NM_003001.2 /DEF=Homo sapiens succinate dehydrogenase complex, subunit C, integral membrane protein, 15kD (SDHC), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=SDHC /PROD=succinate dehydrogenase complex, subunit Cprecursor /DB_XREF=gi:9257243 /UG=Hs.3577 succinate dehydrogenase complex, subunit C, integral membrane protein, 15kD /FL=gb:U57877.1 gb:NM_003001.2	NM_003001		NP_002992
202006_at	1.26E-04	gb:NM_002835.1 /DEF=Homo sapiens protein tyrosine phosphatase, non-receptor type 12 (PTPN12), mRNA. /FEA=mRNA /GEN=PTPN12 /PROD=protein tyrosine phosphatase, non-receptor type12 /DB_XREF=gi:4506286 /UG=Hs.62 protein tyrosine phosphatase, non-receptor type 12 /FL=gb:D13380.1 gb:M93425.1 gb:NM_002835.1	NM_002835		NP_002826

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202009_at	0.016934	gb:NM_007284.1 /DEF=Homo sapiens protein tyrosine kinase 9-like (A6-related protein) (PTK9L), mRNA. /FEA=mRNA /GEN=PTK9L /PROD=protein tyrosine kinase 9-like (A6-related protein) /DB_XREF=gi:6005845 /UG=Hs.6780 protein tyrosine kinase 9-like (A6-related protein) /FL=gb:BC000327.1 gb:BC003161.1 gb:NM_007284.1 gb:AL136773.1	NM_007284		NP_009215
202010_s_at	3.26E-04	gb:NM_021188.1 /DEF=Homo sapiens clones 23667 and 23775 zinc finger protein (LOC57862), mRNA. /FEA=mRNA /GEN=LOC57862 /PROD=clones 23667 and 23775 zinc finger protein /DB_XREF=gi:10863994 /UG=Hs.7137 clones 23667 and 23775 zinc finger protein /FL=gb:NM_021188.1 gb:U90919.1	NM_021188		NP_067011
202018_s_at	0.001679	gb:NM_002343.1 /DEF=Homo sapiens lactotransferrin (LTF), mRNA. /FEA=mRNA /GEN=LTF /PROD=lactotransferrin /DB_XREF=gi:4505042 /UG=Hs.105938 lactotransferrin /FL=gb:AF332168.1 gb:M93150.1 gb:M83202.1 gb:NM_002343.1	NM_002343		NP_002334
202024_at	0.04734	gb:NM_004317.1 /DEF=Homo sapiens arsA (bacterial) arsenite transporter, ATP-binding, homolog 1 (ASNA1), mRNA. /FEA=mRNA /GEN=ASNA1 /PROD=arsA (bacterial) arsenite transporter,ATP-binding, homolog 1 /DB_XREF=gi:4757795 /UG=Hs.165439 arsA (bacterial) arsenite transporter, ATP-binding, homolog 1 /FL=gb:BC002651.1 gb:U60276.1 gb:AF047469.1 gb:NM_004317.1	NM_004317		NP_004308

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202025_x_at	0.00161	gb:NM_001607.2 /DEF=Homo sapiens acetyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Coenzyme A thiolase) (ACAA1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=ACAA1 /PROD=acetyl-Coenzyme A acyltransferase 1 /DB_XREF=gi:6598316 /UG=Hs.166160 acetyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Coenzyme A thiolase) /FL=gb:BC000635.1 gb:NM_001607.2	NM_001607		NP_001598
202026_at	2.05E-04	gb:NM_003002.1 /DEF=Homo sapiens succinate dehydrogenase complex, subunit D, integral membrane protein (SDHD), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=SDHD /PROD=succinate dehydrogenase complex, subunit Dprecursor /DB_XREF=gi:4506864 /UG=Hs.168289 succinate dehydrogenase complex, subunit D, integral membrane protein /FL=gb:BC005263.1 gb:AB006202.1 gb:NM_003002.1	NM_003002		NP_002993
202027_at	0.009682	gb:NM_012264.1 /DEF=Homo sapiens chromosome 22 open reading frame 5 (C22ORF5), mRNA. /FEA=mRNA /GEN=C22ORF5 /PROD=chromosome 22 open reading frame 5 /DB_XREF=gi:7110634 /UG=Hs.182626 chromosome 22 open reading frame 5 /FL=gb:NM_012264.1	NM_012264		NP_036396

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202031_s_at	0.041055	gb:NM_015610.1 /DEF=Homo sapiens DKFZP434J154 protein (DKFZP434J154), mRNA. /FEA=mRNA /GEN=DKFZP434J154 /PROD=DKFZP434J154 protein /DB_XREF=gi:7661579 /UG=Hs.226372 DKFZP434J154 protein /FL=gb:BC004116.1 gb:AL080155.1 gb:NM_015610.1	NM_015610		NP_057087
202032_s_at	0.004341	gb:NM_006122.1 /DEF=Homo sapiens mannosidase, alpha, class 2A, member 2 (MAN2A2), mRNA. /FEA=mRNA /GEN=MAN2A2 /PROD=mannosidase, alpha, class 2A, member 2 /DB_XREF=gi:5540099 /UG=Hs.295605 mannosidase, alpha, class 2A, member 2 /FL=gb:NM_006122.1 gb:L28821.1	NM_006122		NP_006113
202038_at	0.016934	gb:NM_004788.1 /DEF=Homo sapiens ubiquitination factor E4A (homologous to yeast UFD2) (UBE4A), mRNA. /FEA=mRNA /GEN=UBE4A /PROD=ubiquitination factor E4A (homologous to yeastUFD2) /DB_XREF=gi:4759287 /UG=Hs.75275 ubiquitination factor E4A (homologous to yeast UFD2) /FL=gb:D50916.1 gb:NM_004788.1	NM_004788		NP_004779
202040_s_at	1.26E-04	gb:NM_005056.1 /DEF=Homo sapiens retinoblastoma-binding protein 2 (RBBP2), mRNA. /FEA=mRNA /GEN=RBBP2 /PROD=retinoblastoma-binding protein 2 /DB_XREF=gi:4826967 /UG=Hs.76272 retinoblastoma-binding protein 2 /FL=gb:NM_005056.1	NM_005056		NP_005047

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202041_s_at	1.26E-04	gb:NM_004214.3 /DEF=Homo sapiens fibroblast growth factor (acidic) intracellular binding protein (FIBP), mRNA. /FEA=mRNA /GEN=FIBP /PROD=fibroblast growth factor (acidic) intracellular binding protein /DB_XREF=gi:7262377 /UG=Hs.7768 fibroblast growth factor (acidic) intracellular binding protein /FL=gb:AF171944.1 gb:AF171945.1 gb:AF171946.1 gb:AF010187.2 gb:NM_004214.3	NM_004214		NP_004205
202047_s_at	0.00228	chromobox homolog 6	AI458128	Hs.107374	NP_055107
202053_s_at	0.021435	gb:L47162.1 /DEF=Human fatty aldehyde dehydrogenase (FALDH) mRNA, complete cds. /FEA=mRNA /GEN=FALDH /PROD=fatty aldehyde dehydrogenase /DB_XREF=gi:1082035 /UG=Hs.159608 aldehyde dehydrogenase 3 family, member A2 /FL=gb:L47162.1 gb:U46689.1 gb:NM_000382.1	L47162		NP_000373
202054_s_at	0.022752	gb:NM_000382.1 /DEF=Homo sapiens aldehyde dehydrogenase 3 family, member A2 (ALDH3A2), mRNA. /FEA=mRNA /GEN=ALDH3A2 /PROD=aldehyde dehydrogenase 3A2 /DB_XREF=gi:4557302 /UG=Hs.159608 aldehyde dehydrogenase 3 family, member A2 /FL=gb:L47162.1 gb:U46689.1 gb:NM_000382.1	NM_000382		NP_000373
202055_at	7.10E-04	Consensus includes gb:AA652173 /FEA=EST /DB_XREF=gi:2583825 /DB_XREF=est:ns51f05.s1 /CLONE=IMAGE:1187169 /UG=Hs.169149 karyopherin alpha 1 (importin alpha 5) /FL=gb:BC002374.1 gb:BC003009.1 gb:NM_002264.1	NM_002264		NP_002255

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202059_s_at	1.26E-04	gb:NM_002264.1 /DEF=Homo sapiens karyopherin alpha 1 (importin alpha 5) (KPNA1), mRNA. /FEA=mRNA /GEN=KPNA1 /PROD=karyopherin alpha 1 /DB_XREF=gi:4504894 /UG=Hs.169149 karyopherin alpha 1 (importin alpha 5) /FL=gb:BC002374.1 gb:BC003009.1 gb:NM_002264.1	NM_002264		NP_002255
202060_at	0.016934	gb:NM_014633.1 /DEF=Homo sapiens KIAA0155 gene product (KIAA0155), mRNA. /FEA=mRNA /GEN=KIAA0155 /PROD=KIAA0155 gene product /DB_XREF=gi:7661949 /UG=Hs.173288 KIAA0155 gene product /FL=gb:NM_014633.1 gb:D63875.1	NM_014633		NP_055448
202061_s_at	1.26E-04	sel-1 suppressor of lin-12-like (C. elegans)	AI927770	Hs.181300	NP_005056
202066_at	0.010205	Consensus includes gb:AA195259 /FEA=EST /DB_XREF=gi:1784959 /DB_XREF=est:zr36g01.s1 /CLONE=IMAGE:665520 /UG=Hs.183648 protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 1 /FL=gb:NM_003626.1 gb:U22816.1	NM_003626		NP_803172
202068_s_at	0.004341	gb:NM_000527.2 /DEF=Homo sapiens low density lipoprotein receptor (familial hypercholesterolemia) (LDLR), mRNA. /FEA=mRNA /GEN=LDLR /PROD=low density lipoprotein receptor precursor /DB_XREF=gi:8051613 /UG=Hs.213289 low density lipoprotein receptor (familial hypercholesterolemia) /FL=gb:NM_000527.2	NM_000527		NP_000518

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202078_at	0.026891	gb:NM_003653.1 /DEF=Homo sapiens COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 3 (COPS3), mRNA. /FEA=mRNA /GEN=COPS3 /PROD=COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 3 /DB_XREF=gi:4502974 /UG=Hs.6076 COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 3 /FL=gb:BC001891.1 gb:AF031647.1 gb:NM_003653.1 gb:AF098109.1	NM_003653		NP_003644
202079_s_at	0.005339	KIAA1042 protein	AI633774	Hs.6705	NP_055780
202080_s_at	2.75E-04	gb:NM_014965.1 /DEF=Homo sapiens KIAA1042 protein (KIAA1042), mRNA. /FEA=mRNA /GEN=KIAA1042 /PROD=KIAA1042 protein /DB_XREF=gi:7662457 /UG=Hs.6705 KIAA1042 protein /FL=gb:AB028965.1 gb:NM_014965.1	NM_014965		NP_055780
202082_s_at	0.003529	Consensus includes gb:AV748469 /FEA=EST /DB_XREF=gi:10906317 /DB_XREF=est:AV748469 /CLONE=NPCAXA07 /UG=Hs.75232 SEC14 (S. cerevisiae)-like 1 /FL=gb:D67029.1 gb:NM_003003.1	NM_003003		NP_002994
202088_at	0.007779	LIV-1 protein, estrogen regulated	AI635449	Hs.79136	NP_036451
202089_s_at	0.001116	gb:NM_012319.2 /DEF=Homo sapiens LIV-1 protein, estrogen regulated (LIV-1), mRNA. /FEA=mRNA /GEN=LIV-1 /PROD=LIV-1 protein, estrogen regulated /DB_XREF=gi:12751474 /UG=Hs.79136 LIV-1 protein, estrogen regulated /FL=gb:U41060.2 gb:NM_012319.2	NM_012319		NP_036451

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202090_s_at	5.03E-04	gb:NM_006830.1 /DEF=Homo sapiens ubiquinol-cytochrome c reductase (6.4kD) subunit (UQCR), mRNA. /FEA=mRNA /GEN=UQCR /PROD=ubiquinol-cytochrome c reductase (6.4kD)subunit /DB_XREF=gi:5803216 /UG=Hs.8372 ubiquinol-cytochrome c reductase (6.4kD) subunit /FL=gb:D55636.1 gb:NM_006830.1	NM_006830		NP_006821
202092_s_at	0.042536	Consensus includes gb:BF244411 /FEA=EST /DB_XREF=gi:11158342 /DB_XREF=est:601862994F1 /CLONE=IMAGE:4080550 /UG=Hs.9552 binder of Arl Two /FL=gb:BC003087.1 gb:AF126062.1 gb:NM_012106.1	NM_012106		NP_036238
202093_s_at	0.013223	gb:NM_019088.1 /DEF=Homo sapiens hypothetical protein F23149_1 (F23149_1), mRNA. /FEA=mRNA /GEN=F23149_1 /PROD=hypothetical protein F23149_1 /DB_XREF=gi:9506582 /UG=Hs.152894 hypothetical protein F23149_1 /FL=gb:BC000017.1 gb:NM_019088.1	NM_019088		NP_061961
202096_s_at	0.02145	gb:NM_000714.2 /DEF=Homo sapiens benzodiazapine receptor (peripheral) (BZRP), nuclear gene encoding mitochondrial protein, transcript variant PBR, mRNA. /FEA=mRNA /GEN=BZRP /PROD=peripheral benzodiazapine receptor /DB_XREF=gi:6382068 /UG=Hs.202 benzodiazapine receptor (peripheral) /FL=gb:BC001110.1 gb:M36035.1 gb:NM_000714.2	NM_000714		NP_009295

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202097_at	0.001116	gb:NM_005124.1 /DEF=Homo sapiens nucleoporin 153kD (NUP153), mRNA. /FEA=mRNA /GEN=NUP153 /PROD=nucleoporin 153kD /DB_XREF=gi:4826871 /UG=Hs.211608 nucleoporin 153kD /FL=gb:NM_005124.1	NM_005124		NP_005115
202098_s_at	0.007779	gb:NM_001535.1 /DEF=Homo sapiens HMT1 (hnRNP methyltransferase, S. cerevisiae)-like 1 (HRMT1L1), mRNA. /FEA=mRNA /GEN=HRMT1L1 /PROD=HMT1 (hnRNP methyltransferase, S.cerevisiae)-like 1 /DB_XREF=gi:4504494 /UG=Hs.235887 HMT1 (hnRNP methyltransferase, S. cerevisiae)-like 1 /FL=gb:U80213.1 gb:NM_001535.1	NM_001535		NP_001526
202100_at	3.26E-04	v-ral simian leukemia viral oncogene homolog B (ras related; GTP binding protein)	BG169673	Hs.348024	NP_002872
202102_s_at	0.003529	bromodomain containing 4	BF718610	Hs.278675	NP_490597
202105_at	3.26E-04	gb:NM_001551.1 /DEF=Homo sapiens immunoglobulin (CD79A) binding protein 1 (IGBP1), mRNA. /FEA=mRNA /GEN=IGBP1 /PROD=immunoglobulin-binding protein 1 /DB_XREF=gi:4557662 /UG=Hs.3631 immunoglobulin (CD79A) binding protein 1 /FL=gb:BC004137.1 gb:NM_001551.1	NM_001551		NP_001542
202110_at	0.016934	gb:NM_001866.1 /DEF=Homo sapiens cytochrome c oxidase subunit VIIb (COX7B), mRNA. /FEA=mRNA /GEN=COX7B /PROD=cytochrome c oxidase subunit VIIb /DB_XREF=gi:4502990 /UG=Hs.75752 cytochrome c oxidase subunit VIIb /FL=gb:NM_001866.1	NM_001866		NP_001857

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202113_s_at	0.013223	gb:AF043453.1 /DEF=Homo sapiens sorting nexin 2 (SNX2) mRNA, complete cds. /FEA=mRNA /GEN=SNX2 /PROD=sorting nexin 2 /DB_XREF=gi:2827433 /UG=Hs.11183 sorting nexin 2 /FL=gb:BC003382.1 gb:AF043453.1 gb:AF065482.1 gb:NM_003100.1	AF043453		
202117_at	0.041055	Rho GTPase activating protein 1	BG468434	Hs.138860	NP_004299
202118_s_at	0.041055	copine III	AA541758	Hs.14158	NP_003900
202119_s_at	1.26E-04	gb:NM_003909.1 /DEF=Homo sapiens copine III (CPNE3), mRNA. /FEA=mRNA /GEN=CPNE3 /PROD=copine III /DB_XREF=gi:4503014 /UG=Hs.14158 copine III /FL=gb:AB014536.1 gb:NM_003909.1 gb:AF077226.2	NM_003909		NP_003900
202120_x_at	0.010205	gb:NM_004069.2 /DEF=Homo sapiens adaptor-related protein complex 2, sigma 1 subunit (AP2S1), transcript variant AP17, mRNA. /FEA=mRNA /GEN=AP2S1 /PROD=adaptor-related protein complex 2, sigma 1 subunit, isoform AP17 /DB_XREF=gi:11038644 /UG=Hs.119591 adaptor-related protein complex 2, sigma 1 subunit /FL=gb:NM_004069.2	NM_004069		NP_067586
202121_s_at	2.05E-04	gb:NM_014453.1 /DEF=Homo sapiens putative breast adenocarcinoma marker (32kD) (BC-2), mRNA. /FEA=mRNA /GEN=BC-2 /PROD=putative breast adenocarcinoma marker (32kD) /DB_XREF=gi:7656921 /UG=Hs.12107 putative breast adenocarcinoma marker (32kD) /FL=gb:AF042384.1 gb:NM_014453.1	NM_014453		NP_055268

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202122_s_at	0.013223	gb:NM_005817.1 /DEF=Homo sapiens cargo selection protein (mannose 6 phosphate receptor binding protein) (TIP47), mRNA. /FEA=mRNA /GEN=TIP47 /PROD=cargo selection protein (mannose 6 phosphatereceptor binding protein) /DB_XREF=gi:5032182 /UG=Hs.140452 cargo selection protein (mannose 6 phosphate receptor binding protein) /FL=gb:AF057140.1 gb:AF051314.1 gb:AF051315.1 gb:AF055574.1 gb:NM_005817.1	NM_005817		NP_005808
202124_s_at	0.008036	amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3	AV705253	Hs.154248	NP_055864
202125_s_at	0.003171	gb:NM_015049.1 /DEF=Homo sapiens amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3 (ALS2CR3), mRNA. /FEA=mRNA /GEN=ALS2CR3 /PROD=amyotrophic lateral sclerosis 2 (juvenile)chromosome region, candidate 3 /DB_XREF=gi:13027379 /UG=Hs.154248 amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3 /FL=gb:AB038951.1 gb:NM_015049.1	NM_015049		NP_055864
202128_at	5.03E-04	gb:NM_014821.1 /DEF=Homo sapiens KIAA0317 gene product (KIAA0317), mRNA. /FEA=mRNA /GEN=KIAA0317 /PROD=KIAA0317 gene product /DB_XREF=gi:7662051 /UG=Hs.20126 KIAA0317 gene product /FL=gb:AB002315.1 gb:NM_014821.1	NM_014821		NP_055636
202129_s_at	0.001253	sudD suppressor of bimD6 homolog (A. nidulans)	AW006290	Hs.209061	NP_665913
202130_at	1.26E-04	sudD suppressor of bimD6 homolog (A. nidulans)	AW006290	Hs.209061	NP_665913

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202131_s_at	1.26E-04	gb:NM_003831.1 /DEF=Homo sapiens sudD (suppressor of bimD6, Aspergillus nidulans) homolog (SUDD), mRNA. /FEA=mRNA /GEN=SUDD /PROD=sudD (suppressor of bimD6, Aspergillus nidulans)homolog /DB_XREF=gi:4507298 /UG=Hs.209061 sudD (suppressor of bimD6, Aspergillus nidulans) homolog /FL=gb:AF013591.1 gb:NM_003831.1	NM_003831		NP_665913
202135_s_at	0.007779	gb:NM_005735.2 /DEF=Homo sapiens ARP1 (actin-related protein 1, yeast) homolog B (centractin beta) (ACTR1B), mRNA. /FEA=mRNA /GEN=ACTR1B /PROD=actin-related protein 1B /DB_XREF=gi:13325060 /UG=Hs.2477 ARP1 (actin-related protein 1, yeast) homolog B (centractin beta) /FL=gb:NM_005735.2 gb:BC004374.1	NM_005735		NP_005726
202136_at	0.005852	adenovirus 5 E1A binding protein	BE250417	Hs.301449	NP_006615
202140_s_at	0.003171	gb:NM_003992.1 /DEF=Homo sapiens CDC-like kinase 3 (CLK3), transcript variant phclk3, mRNA. /FEA=mRNA /GEN=CLK3 /PROD=CDC-like kinase 3 isoform hclk3 /DB_XREF=gi:4502884 /UG=Hs.73987 CDC-like kinase 3 /FL=gb:BC002555.1 gb:NM_003992.1 gb:L29217.1	NM_003992		NP_003983
202143_s_at	0.020576	gb:NM_006710.1 /DEF=Homo sapiens COP9 homolog (COP9), mRNA. /FEA=mRNA /GEN=COP9 /PROD=COP9 homolog /DB_XREF=gi:5729778 /UG=Hs.75193 COP9 homolog /FL=gb:BC003090.1 gb:U51205.1 gb:NM_006710.1	NM_006710		NP_006701

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202151_s_at	0.038554	gb:NM_016172.1 /DEF=Homo sapiens putative glioblastoma cell differentiation-related (GDBR1), mRNA. /FEA=mRNA /GEN=GDBR1 /PROD=putative glioblastoma celldifferentiation-related protein /DB_XREF=gi:7705380 /UG=Hs.9194 putative glioblastoma cell differentiation-related /FL=gb:BC004967.1 gb:AF176796.1 gb:NM_016172.1	NM_016172		NP_057256
202155_s_at	1.26E-04	gb:NM_005085.1 /DEF=Homo sapiens nucleoporin 214kD (CAIN) (NUP214), mRNA. /FEA=mRNA /GEN=NUP214 /PROD=nucleoporin 214kD (CAIN) /DB_XREF=gi:4826873 /UG=Hs.170285 nucleoporin 214kD (CAIN) /FL=gb:NM_005085.1	NM_005085		NP_705906
202160_at	7.58E-04	gb:NM_004380.1 /DEF=Homo sapiens CREB binding protein (Rubinstein-Taybi syndrome) (CREBBP), mRNA. /FEA=mRNA /GEN=CREBBP /PROD=CREB binding protein /DB_XREF=gi:4758055 /UG=Hs.23598 CREB binding protein (Rubinstein-Taybi syndrome) /FL=gb:U47741.1 gb:NM_004380.1	NM_004380		NP_004371
202162_s_at	3.26E-04	TNFAIP3 interacting protein 1	AI769416	Hs.109281	NP_004770
202163_s_at	0.00161	gb:NM_004779.1 /DEF=Homo sapiens CCR4-NOT transcription complex, subunit 8 (CNOT8), mRNA. /FEA=mRNA /GEN=CNOT8 /PROD=CCR4-NOT transcription complex, subunit 8 /DB_XREF=gi:4758945 /UG=Hs.26703 CCR4-NOT transcription complex, subunit 8 /FL=gb:AF053318.1 gb:NM_004779.1 gb:AL122045.1 gb:AF180476.1	NM_004779		NP_004770

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202164_s_at	1.26E-04	gb:AF180476.1 /DEF=Homo sapiens CALIFp (CALIF) mRNA, complete cds. /FEA=mRNA /GEN=CALIF /PROD=CALIFp /DB_XREF=gi:6856208 /UG=Hs.26703 CCR4-NOT transcription complex, subunit 8 /FL=gb:AF053318.1 gb:NM_004779.1 gb:AL122045.1 gb:AF180476.1	AF180476		NP_004770
202166_s_at	0.041055	gb:NM_006241.1 /DEF=Homo sapiens protein phosphatase 1, regulatory (inhibitor) subunit 2 (PPP1R2), mRNA. /FEA=mRNA /GEN=PPP1R2 /PROD=protein phosphatase 1, regulatory (inhibitor) subunit 2 /DB_XREF=gi:5453945 /UG=Hs.267819 protein phosphatase 1, regulatory (inhibitor) subunit 2 /FL=gb:NM_006241.1	NM_006241		NP_006232
202168_at	0.041055	gb:NM_003187.1 /DEF=Homo sapiens TATA box binding protein (TBP)-associated factor, RNA polymerase II, G, 32kD (TAF2G), mRNA. /FEA=mRNA /GEN=TAF2G /PROD=TATA box binding protein (TBP)-associated factor, RNA polymerase II, G, 32kD /DB_XREF=gi:4507350 /UG=Hs.60679 TATA box binding protein (TBP)-associated factor, RNA polymerase II, G, 32kD /FL=gb:BC003400.1 gb:NM_003187.1 gb:U21858.1 gb:U25112.1 gb:U30504.1	NM_003187		NP_057367
202172_at	0.007779	Consensus includes gb:BG035116 /FEA=EST /DB_XREF=gi:12428927 /DB_XREF=est:602324811F1 /CLONE=IMAGE:4412907 /UG=Hs.6557 zinc finger protein 161 /FL=gb:D28118.1 gb:NM_007146.1	NM_007146		NP_009077

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202181_at	0.00228	gb:NM_014734.1 /DEF=Homo sapiens KIAA0247 gene product (KIAA0247), mRNA. /FEA=mRNA /GEN=KIAA0247 /PROD=KIAA0247 gene product /DB_XREF=gi:7662019 /UG=Hs.82426 KIAA0247 gene product /FL=gb:D87434.1 gb:NM_014734.1	NM_014734		NP_055549
202182_at	0.001116	gb:NM_021078.1 /DEF=Homo sapiens GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 2 (GCN5L2), mRNA. /FEA=mRNA /GEN=GCN5L2 /PROD=GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 2 /DB_XREF=gi:10835100 /UG=Hs.101067 GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 2 /FL=gb:NM_021078.1	NM_021078		NP_066564
202187_s_at	2.05E-04	gb:NM_006243.1 /DEF=Homo sapiens protein phosphatase 2, regulatory subunit B (B56), alpha isoform (PPP2R5A), mRNA. /FEA=mRNA /GEN=PPP2R5A /PROD=protein phosphatase 2, regulatory subunit B(B56), alpha isoform /DB_XREF=gi:5453949 /UG=Hs.155079 protein phosphatase 2, regulatory subunit B (B56), alpha isoform /FL=gb:L42373.1 gb:NM_006243.1	NM_006243		NP_006234
202192_s_at	1.26E-04	gb:NM_005890.1 /DEF=Homo sapiens growth arrest-specific 7 (GAS7), transcript variant b, mRNA. /FEA=mRNA /GEN=GAS7 /PROD=growth arrest-specific 7 isoform b /DB_XREF=gi:5360211 /UG=Hs.226133 growth arrest-specific 7 /FL=gb:AB007854.1 gb:NM_005890.1	NM_005890		NP_005881

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202193_at	0.007779	gb:NM_005569.2 /DEF=Homo sapiens LIM domain kinase 2 (LIMK2), transcript variant 2a, mRNA. /FEA=mRNA /GEN=LIMK2 /PROD=LIM domain kinase 2 isoform 2a /DB_XREF=gi:8051619 /UG=Hs.278027 LIM domain kinase 2 /FL=gb:D45906.1 gb:NM_005569.2	NM_005569		NP_057952
202194_at	0.003171	Consensus includes gb:AL117354 /DEF=Human DNA sequence from clone RP5-976O13 on chromosome 1p21.2-22.2 Contains part of the gene for CGI-100 protein, 3 isoforms of the gene for M96 protein, ESTs, STSs, GSSs and a CpG Island /FEA=mRNA_1 /DB_XREF=gi:6822199 /UG=Hs.296155 CGI-100 protein /FL=gb:AF151858.1 gb:NM_016040.1	AL117354		
202197_at	1.26E-04	gb:NM_021090.1 /DEF=Homo sapiens myotubularin related protein 3 (MTMR3), mRNA. /FEA=mRNA /GEN=MTMR3 /PROD=myotubularin related protein 3 /DB_XREF=gi:10835108 /UG=Hs.63302 myotubularin related protein 3 /FL=gb:NM_021090.1 gb:AB002369.1 gb:AF233438.1	NM_021090		NP_694691
202199_s_at	0.026891	SFRS protein kinase 1	AW082913	Hs.75761	NP_003128
202200_s_at	1.26E-04	gb:NM_003137.1 /DEF=Homo sapiens SFRS protein kinase 1 (SRPK1), mRNA. /FEA=mRNA /GEN=SRPK1 /PROD=SFRS protein kinase 1 /DB_XREF=gi:4507218 /UG=Hs.75761 SFRS protein kinase 1 /FL=gb:NM_003137.1 gb:U09564.1	NM_003137		NP_003128

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202201_at	1.26E-04	gb:NM_000713.1 /DEF=Homo sapiens biliverdin reductase B (flavin reductase (NADPH)) (BLVRB), mRNA. /FEA=mRNA /GEN=BLVRB /PROD=biliverdin reductase B (flavin reductase(NADPH)) /DB_XREF=gi:4502418 /UG=Hs.76289 biliverdin reductase B (flavin reductase (NADPH)) /FL=gb:D26308.1 gb:NM_000713.1	NM_000713		NP_000704
202203_s_at	0.01659	gb:NM_001144.1 /DEF=Homo sapiens autocrine motility factor receptor (AMFR), mRNA. /FEA=mRNA /GEN=AMFR /PROD=autocrine motility factor receptor /DB_XREF=gi:4502074 /UG=Hs.80731 autocrine motility factor receptor /FL=gb:NM_001144.1 gb:L35233.1 gb:AF124145.1	NM_001144		NP_620408
202206_at	2.05E-04	Consensus includes gb:AW450363 /FEA=EST /DB_XREF=gi:6991139 /DB_XREF=est:UI-H-BI3-akn-d-02-0-UI.s1 /CLONE=IMAGE:2734875 /UG=Hs.111554 ADP-ribosylation factor-like 7 /FL=gb:BC001051.1 gb:AB016811.1 gb:NM_005737.2	NM_005737		NP_005728
202207_at	0.00161	Consensus includes gb:BG435404 /FEA=EST /DB_XREF=gi:13341910 /DB_XREF=est:602507678F1 /CLONE=IMAGE:4605066 /UG=Hs.111554 ADP-ribosylation factor-like 7 /FL=gb:BC001051.1 gb:AB016811.1 gb:NM_005737.2	NM_005737		NP_005728
202208_s_at	0.001618	gb:BC001051.1 /DEF=Homo sapiens, ADP-ribosylation factor-like 7, clone MGC:1575, mRNA, complete cds. /FEA=mRNA /PROD=ADP-ribosylation factor-like 7 /DB_XREF=gi:12654450 /UG=Hs.111554 ADP-ribosylation factor-like 7 /FL=gb:BC001051.1 gb:AB016811.1 gb:NM_005737.2	BC001051		NP_005728

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202211_at	3.26E-04	gb:BC005122.1 /DEF=Homo sapiens, ADP-ribosylation factor GTPase activating protein 1, clone MGC:10272, mRNA, complete cds. /FEA=mRNA /PROD=ADP-ribosylation factor GTPase activatingprotein 1 /DB_XREF=gi:13477296 /UG=Hs.13014 ADP-ribosylation factor GTPase activating protein 1 /FL=gb:BC005122.1 gb:AF111847.1 gb:NM_014570.1	BC005122		NP_055385
202214_s_at	0.016934	gb:NM_003588.1 /DEF=Homo sapiens cullin 4B (CUL4B), mRNA. /FEA=mRNA /GEN=CUL4B /PROD=cullin 4B /DB_XREF=gi:13270466 /UG=Hs.155976 cullin 4B /FL=gb:NM_003588.1 gb:AB014595.1	NM_003588		NP_003579
202215_s_at	7.58E-04	gb:NM_014223.2 /DEF=Homo sapiens nuclear transcription factor Y, gamma (NFYC), mRNA. /FEA=mRNA /GEN=NFYC /PROD=nuclear transcription factor Y, gamma /DB_XREF=gi:11496977 /UG=Hs.168157 nuclear transcription factor Y, gamma /FL=gb:NM_014223.2 gb:D85425.1 gb:BC005003.1 gb:D89986.1	NM_014223		NP_055038
202221_s_at	1.26E-04	AV727101 HTC Homo sapiens cDNA clone HTC BGF05 5', mRNA sequence.	AV727101		
202225_at	0.02145	Consensus includes gb:AW612311 /FEA=EST /DB_XREF=gi:7317497 /DB_XREF=est:hg95e07.x1 /CLONE=IMAGE:2953380 /UG=Hs.306088 v-crk avian sarcoma virus CT10 oncogene homolog /FL=gb:D10656.1 gb:NM_016823.1	NM_016823		NP_058431

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202228_s_at	1.26E-04	gb:NM_017455.1 /DEF=Homo sapiens stromal cell derived factor receptor 1 (SDFR1), transcript variant alpha, mRNA. /FEA=mRNA /GEN=SDFR1 /PROD=stromal cell derived factor receptor 1 isoforma /DB_XREF=gi:9257239 /UG=Hs.6354 stromal cell derived factor receptor 1 /FL=gb:AF109127.1 gb:NM_017455.1	NM_017455		NP_059429
202230_s_at	0.016934	gb:NM_006387.2 /DEF=Homo sapiens protein with polyglutamine repeat; calcium (ca2+) homeostasis endoplasmic reticulum protein (ERPROT213-21), mRNA. /FEA=mRNA /GEN=ERPROT213-21 /PROD=protein with polyglutamine repeat; calcium(ca2+) homeostasis endoplasmic reticulum protein /DB_XREF=gi:11055968 /UG=Hs.6430 protein with polyglutamine repeat; calcium (ca2+) homeostasis endoplasmic reticulum protein /FL=gb:U94836.2 gb:NM_006387.2	NM_006387		NP_006378
202233_s_at	0.010205	gb:NM_006004.1 /DEF=Homo sapiens ubiquinol-cytochrome c reductase hinge protein (UQCRH), mRNA. /FEA=mRNA /GEN=UQCRH /PROD=ubiquinol-cytochrome c reductase hinge protein /DB_XREF=gi:5174744 /UG=Hs.73818 ubiquinol-cytochrome c reductase hinge protein /FL=gb:BC001426.1 gb:BC001934.1 gb:M36647.1 gb:NM_006004.1	NM_006004		NP_005995

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202236_s_at	0.026891	gb:NM_003051.1 /DEF=Homo sapiens solute carrier family 16 (monocarboxylic acid transporters), member 1 (SLC16A1), mRNA. /FEA=mRNA /GEN=SLC16A1 /PROD=solute carrier family 16 (monocarboxylic acid transporters), member 1 /DB_XREF=gi:4506982 /UG=Hs.75231 solute carrier family 16 (monocarboxylic acid transporters), member 1 /FL=gb:NM_003051.1 gb:L31801.1	NM_003051		NP_003042
202239_at	7.58E-04	gb:NM_006437.2 /DEF=Homo sapiens ADP-ribosyltransferase (NAD ⁺ ; poly (ADP-ribose) polymerase)-like 1 (ADPRTL1), mRNA. /FEA=mRNA /GEN=ADPRTL1 /PROD=poly(ADP-ribosyl)transferase-like 1 /DB_XREF=gi:11496990 /UG=Hs.77225 ADP-ribosyltransferase (NAD ⁺ ; poly (ADP-ribose) polymerase)-like 1 /FL=gb:NM_006437.2 gb:AF057160.1 gb:AF158255.1	NM_006437		NP_006428
202241_at	0.00228	gb:NM_025195.1 /DEF=Homo sapiens phosphoprotein regulated by mitogenic pathways (C8FW), mRNA. /FEA=mRNA /GEN=C8FW /PROD=G-protein-coupled receptor induced protein /DB_XREF=gi:13399327 /UG=Hs.7837 phosphoprotein regulated by mitogenic pathways /FL=gb:AF205437.1 gb:NM_025195.1	NM_025195		NP_079471

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202243_s_at	2.05E-04	gb:NM_002796.1 /DEF=Homo sapiens proteasome (prosome, macropain) subunit, beta type, 4 (PSMB4), mRNA. /FEA=mRNA /GEN=PSMB4 /PROD=proteasome (prosome, macropain) subunit, betatype, 4 /DB_XREF=gi:4506198 /UG=Hs.89545 proteasome (prosome, macropain) subunit, beta type, 4 /FL=gb:BC000331.1 gb:NM_002796.1 gb:D26600.1	NM_002796		NP_002787
202244_at	0.005852	gb:NM_002796.1 /DEF=Homo sapiens proteasome (prosome, macropain) subunit, beta type, 4 (PSMB4), mRNA. /FEA=mRNA /GEN=PSMB4 /PROD=proteasome (prosome, macropain) subunit, betatype, 4 /DB_XREF=gi:4506198 /UG=Hs.89545 proteasome (prosome, macropain) subunit, beta type, 4 /FL=gb:BC000331.1 gb:NM_002796.1 gb:D26600.1	NM_002796		NP_002787
202247_s_at	7.58E-04	metastasis associated 1	BE561596	Hs.101448	NP_004680
202252_at	1.92E-04	gb:NM_002870.1 /DEF=Homo sapiens RAB13, member RAS oncogene family (RAB13), mRNA. /FEA=mRNA /GEN=RAB13 /PROD=RAB13, member RAS oncogene family /DB_XREF=gi:4506362 /UG=Hs.151536 RAB13, member RAS oncogene family /FL=gb:BC000799.1 gb:NM_002870.1	NM_002870		NP_002861

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202257_s_at	0.001713	gb:NM_006110.1 /DEF=Homo sapiens CD2 antigen (cytoplasmic tail)-binding protein 2 (CD2BP2), mRNA. /FEA=mRNA /GEN=CD2BP2 /PROD=CD2 antigen (cytoplasmic tail)-binding protein2 /DB_XREF=gi:5174408 /UG=Hs.202677 CD2 antigen (cytoplasmic tail)-binding protein 2 /FL=gb:BC000495.1 gb:BC001947.1 gb:AF104222.1 gb:NM_006110.1	NM_006110		NP_006101
202263_at	1.26E-04	gb:NM_016243.1 /DEF=Homo sapiens cytochrome b5 reductase 1 (B5R.1) (LOC51706), mRNA. /FEA=mRNA /GEN=LOC51706 /PROD=cytochrome b5 reductase 1 (B5R.1) /DB_XREF=gi:7706454 /UG=Hs.289113 cytochrome b5 reductase 1 (B5R.1) /FL=gb:AF169481.1 gb:AF125533.1 gb:NM_016243.1	NM_016243		NP_057327
202269_x_at	3.26E-04	gb:BC002666.1 /DEF=Homo sapiens, guanylate binding protein 1, interferon-inducible, 67kD, clone MGC:3949, mRNA, complete cds. /FEA=mRNA /PROD=guanylate binding protein 1,interferon-inducible, 67kD /DB_XREF=gi:12803662 /UG=Hs.62661 guanylate binding protein 1, interferon-inducible, 67kD /FL=gb:BC002666.1 gb:M55542.1 gb:NM_002053.1	BC002666		NP_002044
202275_at	0.026891	gb:NM_000402.1 /DEF=Homo sapiens glucose-6-phosphate dehydrogenase (G6PD), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=G6PD /PROD=glucose-6-phosphate dehydrogenase /DB_XREF=gi:4503844 /UG=Hs.80206 glucose-6-phosphate dehydrogenase /FL=gb:BC000337.1 gb:M21248.1 gb:NM_000402.1	NM_000402		NP_000393

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202277_at	0.001116	serine palmitoyltransferase, long chain base subunit 1	AL568804	Hs.90458	
202278_s_at	0.004659	gb:NM_006415.1 /DEF=Homo sapiens serine palmitoyltransferase, long chain base subunit 1 (SPTLC1), mRNA. /FEA=mRNA /GEN=SPTLC1 /PROD=serine palmitoyltransferase, long chain basesubunit 1 /DB_XREF=gi:5454083 /UG=Hs.90458 serine palmitoyltransferase, long chain base subunit 1 /FL=gb:NM_006415.1	NM_006415		NP_847894
202279_at	0.016934	gb:NM_004894.1 /DEF=Homo sapiens chromosome 14 open reading frame 2 (C14ORF2), mRNA. /FEA=mRNA /GEN=C14ORF2 /PROD=chromosome 14 open reading frame 2 /DB_XREF=gi:4758939 /UG=Hs.109052 chromosome 14 open reading frame 2 /FL=gb:BC000429.1 gb:BC001944.1 gb:AF054175.1 gb:NM_004894.1	NM_004894		NP_004885
202281_at	0.005039	gb:NM_005255.1 /DEF=Homo sapiens cyclin G associated kinase (GAK), mRNA. /FEA=mRNA /GEN=GAK /PROD=cyclin G associated kinase /DB_XREF=gi:4885250 /UG=Hs.153227 cyclin G associated kinase /FL=gb:D88435.1 gb:NM_005255.1	NM_005255		NP_005246

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202284_s_at	0.013575	gb:NM_000389.1 /DEF=Homo sapiens cyclin-dependent kinase inhibitor 1A (p21, Cip1) (CDKN1A), mRNA. /FEA=mRNA /GEN=CDKN1A /PROD=cyclin-dependent kinase inhibitor 1A (p21,Cip1) /DB_XREF=gi:11386202 /UG=Hs.179665 cyclin-dependent kinase inhibitor 1A (p21, Cip1) /FL=gb:NM_000389.1 gb:BC000275.1 gb:BC001935.1 gb:U03106.1 gb:L26165.1 gb:L25610.1 gb:U09579.1	NM_000389		NP_510867
202286_s_at	5.67E-04	GA733-1 protein precursor; Human gastrointestinal tumor-associated antigen GA733-1 protein gene, complete cds, clone 05516.	NM_002353		NP_002344
202290_at	0.003529	gb:NM_014891.1 /DEF=Homo sapiens PDGF associated protein (HASPP28), mRNA. /FEA=mRNA /GEN=HASPP28 /PROD=PDGF associated protein /DB_XREF=gi:7657440 /UG=Hs.278426 PDGFA associated protein 1 /FL=gb:U41745.1 gb:BC000684.1 gb:NM_014891.1	NM_014891		NP_055706
202293_at	0.004205	Consensus includes gb:AW168948 /FEA=EST /DB_XREF=gi:6400473 /DB_XREF=est:xj15f07.x1 /CLONE=IMAGE:2657317 /UG=Hs.286148 stromal antigen 1 /FL=gb:NM_005862.1	NM_005862		NP_005853
202295_s_at	1.26E-04	gb:NM_004390.1 /DEF=Homo sapiens cathepsin H (CTSH), mRNA. /FEA=mRNA /GEN=CTSH /PROD=cathepsin H /DB_XREF=gi:4758095 /UG=Hs.288181 cathepsin H /FL=gb:BC002479.1 gb:NM_004390.1	NM_004390		NP_683880

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202297_s_at	3.26E-04	gb:AF157324.1 /DEF=Homo sapiens RER1 protein (RER1) mRNA, complete cds. /FEA=mRNA /GEN=RER1 /PROD=RER1 protein /DB_XREF=gi:7688698 /UG=Hs.40500 similar to S. cerevisiae RER1 /FL=gb:BC004965.1 gb:NM_007033.1 gb:AF157324.1	AF157324		NP_008964
202298_at	1.26E-04	gb:NM_004541.2 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1 (7.5kD, MWFE) (NDUFA1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=NDUFA1 /PROD=NADH dehydrogenase (ubiquinone) 1 alphasubcomplex, 1 /DB_XREF=gi:13699820 /UG=Hs.74823 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1 (7.5kD, MWFE) /FL=gb:BC000266.1 gb:NM_004541.2 gb:U54993.1	NM_004541		NP_004532
202299_s_at	1.26E-04	gb:NM_006402.1 /DEF=Homo sapiens hepatitis B virus x-interacting protein (9.6kD) (XIP), mRNA. /FEA=mRNA /GEN=XIP /PROD=hepatitis B virus x-interacting protein /DB_XREF=gi:5454169 /UG=Hs.80464 hepatitis B virus x-interacting protein (9.6kD) /FL=gb:AF029890.1 gb:NM_006402.1	NM_006402		NP_006393
202300_at	1.26E-04	gb:NM_006402.1 /DEF=Homo sapiens hepatitis B virus x-interacting protein (9.6kD) (XIP), mRNA. /FEA=mRNA /GEN=XIP /PROD=hepatitis B virus x-interacting protein /DB_XREF=gi:5454169 /UG=Hs.80464 hepatitis B virus x-interacting protein (9.6kD) /FL=gb:AF029890.1 gb:NM_006402.1	NM_006402		NP_006393

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202303_x_at	0.041055	gb:NM_003601.1 /DEF=Homo sapiens SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5 (SMARCA5), mRNA. /FEA=mRNA /GEN=SMARCA5 /PROD=SWISNF related, matrix associated, actindependent regulator of chromatin, subfamily a, member 5 /DB_XREF=gi:4507074 /UG=Hs.9456 SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5 /FL=gb:AB010882.1 gb:NM_003601.1	NM_003601		NP_003592
202304_at	5.03E-04	gb:NM_014923.1 /DEF=Homo sapiens KIAA0970 protein (KIAA0970), mRNA. /FEA=mRNA /GEN=KIAA0970 /PROD=KIAA0970 protein /DB_XREF=gi:7662419 /UG=Hs.103329 KIAA0970 protein /FL=gb:AB023187.1 gb:NM_014923.1	NM_014923		
202313_at	2.05E-04	gb:NM_002717.1 /DEF=Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform (PPP2R2A), mRNA. /FEA=mRNA /GEN=PPP2R2A /PROD=protein phosphatase 2 (formerly 2A), regulatorysubunit B (PR 52), alpha isoform /DB_XREF=gi:4506018 /UG=Hs.179574 protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform /FL=gb:M64929.1 gb:NM_002717.1	NM_002717		NP_002708

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202317_s_at	0.013223	gb:NM_006048.1 /DEF=Homo sapiens ubiquitination factor E4B (homologous to yeast UFD2) (UBE4B), mRNA. /FEA=mRNA /GEN=UBE4B /PROD=ubiquitination factor E4B (homologous to yeastUFD2) /DB_XREF=gi:5174482 /UG=Hs.24594 ubiquitination factor E4B (homologous to yeast UFD2) /FL=gb:AF043117.1 gb:NM_006048.1	NM_006048		NP_006039
202318_s_at	0.016934	gb:AF306508.1 /DEF=Homo sapiens SUMO-1 specific protease FKSG6 mRNA, complete cds. /FEA=mRNA /PROD=SUMO-1 specific protease FKSG6 /DB_XREF=gi:11096243 /UG=Hs.27197 SUMO-1-specific protease /FL=gb:AF307849.1 gb:AF306508.1 gb:AF196304.1 gb:NM_015571.1	AF306508		NP_056386
202325_s_at	7.58E-04	gb:NM_001685.1 /DEF=Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit F6 (ATP5J), mRNA. /FEA=mRNA /GEN=ATP5J /PROD=ATP synthase, H ⁺ transporting, mitochondrial /DB_XREF=gi:4502292 /UG=Hs.73851 ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit F6 /FL=gb:M37104.1 gb:M73031.1 gb:NM_001685.1 gb:AL110183.1	NM_001685		NP_001676
202329_at	0.004341	gb:NM_004383.1 /DEF=Homo sapiens c-src tyrosine kinase (CSK), mRNA. /FEA=mRNA /GEN=CSK /PROD=c-src tyrosine kinase /DB_XREF=gi:4758077 /UG=Hs.77793 c-src tyrosine kinase /FL=gb:NM_004383.1	NM_004383		NP_004374

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202330_s_at	0.00884	gb:NM_003362.1 /DEF=Homo sapiens uracil-DNA glycosylase (UNG), mRNA. /FEA=mRNA /GEN=UNG /PROD=uracil-DNA glycosylase /DB_XREF=gi:6224978 /UG=Hs.78853 uracil-DNA glycosylase /FL=gb:NM_003362.1	NM_003362		NP_550433
202333_s_at	0.003171	ubiquitin-conjugating enzyme E2B (RAD6 homolog)	AA877765	Hs.811	NP_003328
202334_s_at	0.001116	ubiquitin-conjugating enzyme E2B (RAD6 homolog)	AA877765	Hs.811	NP_003328
202336_s_at	0.02145	gb:NM_000919.1 /DEF=Homo sapiens peptidylglycine alpha-amidating monooxygenase (PAM), mRNA. /FEA=mRNA /GEN=PAM /PROD=peptidylglycine alpha-amidating monooxygenase /DB_XREF=gi:4505602 /UG=Hs.83920 peptidylglycine alpha-amidating monooxygenase /FL=gb:M37721.1 gb:NM_000919.1	NM_000919		NP_620177
202343_x_at	3.26E-04	gb:NM_001862.1 /DEF=Homo sapiens cytochrome c oxidase subunit Vb (COX5B), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=COX5B /PROD=cytochrome c oxidase subunit Vb /DB_XREF=gi:4502982 /UG=Hs.1342 cytochrome c oxidase subunit Vb /FL=gb:M19961.1 gb:NM_001862.1	NM_001862		NP_001853
202346_at	1.26E-04	gb:NM_005339.2 /DEF=Homo sapiens huntingtin interacting protein 2 (HIP2), mRNA. /FEA=mRNA /GEN=HIP2 /PROD=huntingtin interacting protein 2 /DB_XREF=gi:12545382 /UG=Hs.155485 huntingtin interacting protein 2 /FL=gb:NM_005339.2 gb:U58522.1 gb:AB022435.1	NM_005339		NP_005330

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202347_s_at	5.03E-04	gb:AB022435.1 /DEF=Homo sapiens LIG mRNA for E2 ubiquitin-conjugating enzyme, complete cds. /FEA=mRNA /GEN=LIG /PROD=E2 ubiquitin-conjugating enzyme /DB_XREF=gi:4996607 /UG=Hs.155485 huntingtin interacting protein 2 /FL=gb:NM_005339.2 gb:U58522.1 gb:AB022435.1	AB022435		NP_005330
202348_s_at	0.033381	gb:BC000674.1 /DEF=Homo sapiens, dystonia 1, torsion (autosomal dominant; torsin A), clone MGC:1558, mRNA, complete cds. /FEA=mRNA /PROD=dystonia 1, torsion (autosomal dominant; torsinA) /DB_XREF=gi:12653776 /UG=Hs.19261 dystonia 1, torsion (autosomal dominant; torsin A) /FL=gb:BC000674.1 gb:AF007871.1 gb:NM_000113.1	BC000674		NP_000104
202349_at	1.32E-04	gb:NM_000113.1 /DEF=Homo sapiens dystonia 1, torsion (autosomal dominant; torsin A) (DYT1), mRNA. /FEA=mRNA /GEN=DYT1 /PROD=torsin A /DB_XREF=gi:4557540 /UG=Hs.19261 dystonia 1, torsion (autosomal dominant; torsin A) /FL=gb:BC000674.1 gb:AF007871.1 gb:NM_000113.1	NM_000113		NP_000104
202352_s_at	1.26E-04	proteasome (prosome, macropain) 26S subunit, non-ATPase, 12	AI446530	Hs.4295	NP_777360
202358_s_at	0.003166	KIAA0254 gene product	BG434168	Hs.76906	NP_055573
202359_s_at	0.0375	gb:NM_014758.1 /DEF=Homo sapiens KIAA0254 gene product (KIAA0254), mRNA. /FEA=mRNA /GEN=KIAA0254 /PROD=KIAA0254 gene product /DB_XREF=gi:7662025 /UG=Hs.76906 KIAA0254 gene product /FL=gb:D87443.1 gb:NM_014758.1	NM_014758		NP_055573

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202360_at	1.26E-04	gb:NM_014757.2 /DEF=Homo sapiens mastermind (Drosophila), homolog of (MAML1), mRNA. /FEA=mRNA /GEN=MAML1 /PROD=homolog of Drosophila mastermind /DB_XREF=gi:13376996 /UG=Hs.76986 mastermind (Drosophila), homolog of /FL=gb:NM_014757.2 gb:D83785.1	NM_014757		NP_055572
202362_at	7.58E-04	gb:NM_002884.1 /DEF=Homo sapiens RAP1A, member of RAS oncogene family (RAP1A), mRNA. /FEA=mRNA /GEN=RAP1A /PROD=RAP1A, member of RAS oncogene family /DB_XREF=gi:4506412 /UG=Hs.865 RAP1A, member of RAS oncogene family /FL=gb:NM_002884.1 gb:M22995.1	NM_002884		NP_002875
202364_at	3.26E-04	gb:NM_005962.1 /DEF=Homo sapiens MAX-interacting protein 1 (MXI1), mRNA. /FEA=mRNA /GEN=MXI1 /PROD=MAX-interacting protein 1 /DB_XREF=gi:5174596 /UG=Hs.118630 MAX-interacting protein 1 /FL=gb:L07648.1 gb:NM_005962.1 gb:D63940.1	NM_005962		NP_569157
202367_at	8.68E-04	gb:NM_001913.1 /DEF=Homo sapiens cut (Drosophila)-like 1 (CCAAT displacement protein) (CUTL1), mRNA. /FEA=mRNA /GEN=CUTL1 /PROD=cut (Drosophila)-like 1 (CCAAT displacementprotein) /DB_XREF=gi:4503168 /UG=Hs.147049 cut (Drosophila)-like 1 (CCAAT displacement protein) /FL=gb:NM_001913.1 gb:L12579.1	NM_001913		NP_001904

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202371_at	0.006724	gb:NM_024863.1 /DEF=Homo sapiens hypothetical protein FLJ21174 (FLJ21174), mRNA. /FEA=mRNA /GEN=FLJ21174 /PROD=hypothetical protein FLJ21174 /DB_XREF=gi:13376293 /UG=Hs.194329 hypothetical protein FLJ21174 /FL=gb:AF271783.1 gb:NM_024863.1	NM_024863		NP_079139
202373_s_at	0.00161	gb:AF255648.1 /DEF=Homo sapiens rGAP-iso mRNA, complete cds. /FEA=mRNA /PROD=rGAP-iso /DB_XREF=gi:12005820 /UG=Hs.197289 rab3 GTPase-activating protein, non-catalytic subunit (150kD) /FL=gb:AF255648.1 gb:AF004828.1 gb:NM_012414.1	AF255648		NP_036546
202374_s_at	6.79E-04	gb:NM_012414.1 /DEF=Homo sapiens rab3 GTPase-activating protein, non-catalytic subunit (150kD) (RAB3-GAP150), mRNA. /FEA=mRNA /GEN=RAB3-GAP150 /PROD=rab3 GTPase-activating protein, non-catalytic subunit (150kD) /DB_XREF=gi:6912619 /UG=Hs.197289 rab3 GTPase-activating protein, non-catalytic subunit (150kD) /FL=gb:AF255648.1 gb:AF004828.1 gb:NM_012414.1	NM_012414		NP_036546
202375_at	1.26E-04	gb:NM_014822.1 /DEF=Homo sapiens SEC24 (S. cerevisiae) related gene family, member D (SEC24D), mRNA. /FEA=mRNA /GEN=SEC24D /PROD=SEC24 (S. cerevisiae) related gene family, member D /DB_XREF=gi:7662658 /UG=Hs.19822 SEC24 (S. cerevisiae) related gene family, member D /FL=gb:AB018298.1 gb:AF130464.2 gb:NM_014822.1	NM_014822		NP_055637
202377_at	1.26E-04	ESTs	AW026535	Hs.23581	NP_059996
202379_s_at	1.26E-04	natural killer-tumor recognition sequence	AI361805	Hs.241493	NP_005376

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202381_at	1.26E-04	gb:NM_003816.1 /DEF=Homo sapiens a disintegrin and metalloproteinase domain 9 (meltrin gamma) (ADAM9), mRNA. /FEA=mRNA /GEN=ADAM9 /PROD=a disintegrin and metalloproteinase domain 9preproprotein /DB_XREF=gi:4501914 /UG=Hs.2442 a disintegrin and metalloproteinase domain 9 (meltrin gamma) /FL=gb:U41766.1 gb:NM_003816.1	NM_003816		NP_003807
202382_s_at	8.68E-04	gb:NM_005471.1 /DEF=Homo sapiens glucosamine-6-phosphate isomerase (GNPI), mRNA. /FEA=mRNA /GEN=GNPI /PROD=glucosamine-6-phosphate isomerase /DB_XREF=gi:13027377 /UG=Hs.278500 glucosamine-6-phosphate isomerase /FL=gb:NM_005471.1 gb:AF029914.1 gb:AF048826.1 gb:D31766.1	NM_005471		NP_005462
202386_s_at	1.26E-04	gb:NM_019081.1 /DEF=Homo sapiens KIAA0430 gene product (KIAA0430), mRNA. /FEA=mRNA /GEN=KIAA0430 /PROD=hypothetical protein A-362G6.1 /DB_XREF=gi:11464998 /UG=Hs.30909 KIAA0430 gene product /FL=gb:NM_019081.1	NM_019081		NP_061954
202387_at	1.26E-04	gb:NM_004323.2 /DEF=Homo sapiens BCL2-associated athanogene (BAG1), mRNA. /FEA=mRNA /GEN=BAG1 /PROD=BCL2-associated athanogene /DB_XREF=gi:7549801 /UG=Hs.41714 BCL2-associated athanogene /FL=gb:BC001936.1 gb:AF022224.1 gb:U46917.1 gb:NM_004323.2	NM_004323		NP_004314

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202391_at	0.02145	gb:NM_006317.1 /DEF=Homo sapiens brain acid-soluble protein 1 (BASP1), mRNA. /FEA=mRNA /GEN=BASP1 /PROD=brain acid-soluble protein 1 /DB_XREF=gi:5453749 /UG=Hs.79516 brain abundant, membrane attached signal protein 1 /FL=gb:BC000518.1 gb:AF039656.1 gb:NM_006317.1	NM_006317		NP_006308
202393_s_at	7.58E-04	gb:NM_005655.1 /DEF=Homo sapiens TGFB inducible early growth response (TIEG), mRNA. /FEA=mRNA /GEN=TIEG /PROD=TGFB inducible early growth response /DB_XREF=gi:5032176 /UG=Hs.82173 TGFB inducible early growth response /FL=gb:U21847.1 gb:NM_005655.1	NM_005655		NP_005646
202395_at	1.26E-04	gb:NM_006178.1 /DEF=Homo sapiens N-ethylmaleimide-sensitive factor (NSF), mRNA. /FEA=mRNA /GEN=NSF /PROD=N-ethylmaleimide-sensitive factor /DB_XREF=gi:11079227 /UG=Hs.108802 N-ethylmaleimide-sensitive factor /FL=gb:AF102846.2 gb:NM_006178.1 gb:AF135168.1	NM_006178		NP_006169
202397_at	0.004341	gb:NM_005796.1 /DEF=Homo sapiens nuclear transport factor 2 (placental protein 15) (PP15), mRNA. /FEA=mRNA /GEN=PP15 /PROD=nuclear transport factor 2 (placental protein15) /DB_XREF=gi:5031984 /UG=Hs.151734 nuclear transport factor 2 (placental protein 15) /FL=gb:U43939.1 gb:BC002348.1 gb:NM_005796.1	NM_005796		NP_005787

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202398_at	0.005692	gb:BC002785.1 /DEF=Homo sapiens, adaptor-related protein complex 3, sigma 2 subunit, clone MGC:3577, mRNA, complete cds. /FEA=mRNA /PROD=adaptor-related protein complex 3, sigma 2subunit /DB_XREF=gi:12803880 /UG=Hs.154782 adaptor-related protein complex 3, sigma 2 subunit /FL=gb:BC002785.1 gb:NM_005829.1	BC002785		NP_005820
202399_s_at	1.26E-04	gb:NM_005829.1 /DEF=Homo sapiens adaptor-related protein complex 3, sigma 2 subunit (AP3S2), mRNA. /FEA=mRNA /GEN=AP3S2 /PROD=adaptor-related protein complex 3, sigma 2subunit /DB_XREF=gi:5031580 /UG=Hs.154782 adaptor-related protein complex 3, sigma 2 subunit /FL=gb:BC002785.1 gb:NM_005829.1	NM_005829		NP_005820
202401_s_at	7.57E-04	gb:NM_003131.1 /DEF=Homo sapiens serum response factor (c-fos serum response element-binding transcription factor) (SRF), mRNA. /FEA=mRNA /GEN=SRF /PROD=serum response factor (c-fos serum responseelement-binding transcription factor) /DB_XREF=gi:4507204 /UG=Hs.155321 serum response factor (c-fos serum response element-binding transcription factor) /FL=gb:J03161.1 gb:NM_003131.1	NM_003131		NP_003122
202402_s_at	0.00228	gb:NM_001751.1 /DEF=Homo sapiens cysteinyl-tRNA synthetase (CARS), mRNA. /FEA=mRNA /GEN=CARS /PROD=cysteinyl-tRNA synthetase /DB_XREF=gi:10835050 /UG=Hs.159604 cysteinyl-tRNA synthetase /FL=gb:NM_001751.1 gb:BC002880.1 gb:AF288206.1 gb:AF288207.1	NM_001751		NP_644802

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202412_s_at	0.007052	ubiquitin specific protease 1	AW499935	Hs.35086	NP_003359
202413_s_at	0.02145	gb:NM_003368.1 /DEF=Homo sapiens ubiquitin specific protease 1 (USP1), mRNA. /FEA=mRNA /GEN=USP1 /PROD=ubiquitin specific protease 1 /DB_XREF=gi:4507850 /UG=Hs.35086 ubiquitin specific protease 1 /FL=gb:AB014458.1 gb:AF117386.1 gb:NM_003368.1 gb:AL117575.1	NM_003368		NP_003359
202416_at	0.035171	gb:NM_003315.1 /DEF=Homo sapiens tetratricopeptide repeat domain 2 (TTC2), mRNA. /FEA=mRNA /GEN=TTC2 /PROD=tetratricopeptide repeat domain 2 /DB_XREF=gi:4507712 /UG=Hs.5542 DnaJ (Hsp40) homolog, subfamily C, member 7 /FL=gb:U46571.1 gb:NM_003315.1	NM_003315		NP_003306
202420_s_at	0.013223	gb:NM_001357.2 /DEF=Homo sapiens DEADH (Asp-Glu-Ala-AspHis) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophysin) (DDX9), transcript variant 1, mRNA. /FEA=mRNA /GEN=DDX9 /PROD=DEADH (Asp-Glu-Ala-AspHis) box polypeptide 9, isoform 1 /DB_XREF=gi:13514819 /UG=Hs.74578 DEADH (Asp-Glu-Ala-AspHis) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophysin) /FL=gb:NM_001357.2 gb:L13848.1	NM_001357		NP_085077
202422_s_at	0.004341	gb:NM_022977.1 /DEF=Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 4 (FACL4), transcript variant 2, mRNA. /FEA=mRNA /GEN=FACL4 /PROD=long-chain fatty-acid-Coenzyme A ligase 4, isoform 2 /DB_XREF=gi:12669908 /UG=Hs.81452 fatty-acid-Coenzyme A ligase, long-chain 4 /FL=gb:NM_022977.1	NM_022977		NP_075266

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202423_at	0.004341	gb:NM_006766.1 /DEF=Homo sapiens zinc finger protein 220 (ZNF220), mRNA. /FEA=mRNA /GEN=ZNF220 /PROD=zinc finger protein 220 /DB_XREF=gi:5803097 /UG=Hs.82210 zinc finger protein 220 /FL=gb:U47742.1 gb:NM_006766.1	NM_006766		NP_006757
202427_s_at	3.26E-04	gb:NM_015415.1 /DEF=Homo sapiens DKFZP564B167 protein (DKFZP564B167), mRNA. /FEA=mRNA /GEN=DKFZP564B167 /PROD=DKFZP564B167 protein /DB_XREF=gi:7661601 /UG=Hs.76285 DKFZP564B167 protein /FL=gb:AL110297.1 gb:NM_015415.1	NM_015415		NP_056230
202429_s_at	0.00228	gb:AL353950.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761L0516 (from clone DKFZp761L0516); complete cds. /FEA=mRNA /GEN=DKFZp761L0516 /PROD=hypothetical protein /DB_XREF=gi:7669991 /UG=Hs.272458 protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha) /FL=gb:J05480.1 gb:L14778.1 gb:NM_000944.1 gb:AL353950.1	AL353950		NP_000935
202430_s_at	5.03E-04	gb:NM_021105.1 /DEF=Homo sapiens phospholipid scramblase 1 (PLSCR1), mRNA. /FEA=mRNA /GEN=PLSCR1 /PROD=phospholipid scramblase 1 /DB_XREF=gi:10863876 /UG=Hs.198282 phospholipid scramblase 1 /FL=gb:NM_021105.1 gb:AB006746.1 gb:AF098642.1	NM_021105		NP_066928

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202432_at	3.26E-04	gb:NM_021132.1 /DEF=Homo sapiens protein phosphatase 3 (formerly 2B), catalytic subunit, beta isoform (calcineurin A beta) (PPP3CB), mRNA. /FEA=mRNA /GEN=PPP3CB /PROD=protein phosphatase 3 (formerly 2B), catalyticsubunit, beta isoform (calcineurin A beta) /DB_XREF=gi:11036639 /UG=Hs.151531 protein phosphatase 3 (formerly 2B), catalytic subunit, beta isoform (calcineurin A beta) /FL=gb:NM_021132.1 gb:M29551.1	NM_021132		NP_066955
202433_at	1.26E-04	gb:NM_005827.1 /DEF=Homo sapiens UDP-galactose transporter related (UGTREL1), mRNA. /FEA=mRNA /GEN=UGTREL1 /PROD=UDP-galactose transporter related /DB_XREF=gi:5032212 /UG=Hs.154073 UDP-galactose transporter related /FL=gb:D87989.1 gb:NM_005827.1	NM_005827		NP_005818
202437_s_at	0.005852	gb:NM_000104.2 /DEF=Homo sapiens cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile) (CYP1B1), mRNA. /FEA=mRNA /GEN=CYP1B1 /PROD=cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 /DB_XREF=gi:13325059 /UG=Hs.154654 cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile) /FL=gb:NM_000104.2 gb:U03688.1	NM_000104		NP_000095
202441_at	1.26E-04	similar to Caenorhabditis elegans protein C42C1.9	AL568449	Hs.285818	NP_006450

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202442_at	1.26E-04	gb:NM_001284.1 /DEF=Homo sapiens adaptor-related protein complex 3, sigma 1 subunit (AP3S1), mRNA. /FEA=mRNA /GEN=AP3S1 /PROD=adaptor-related protein complex 3, sigma 1 subunit /DB_XREF=gi:4502860 /UG=Hs.80917 adaptor-related protein complex 3, sigma 1 subunit /FL=gb:BC000804.1 gb:D63643.1 gb:U91932.1 gb:NM_001284.1	NM_001284		NP_001275
202443_x_at	7.58E-04	Notch homolog 2 (Drosophila)	AA291203	Hs.131286	NP_077719
202444_s_at	0.045257	gb:NM_006459.1 /DEF=Homo sapiens similar to Caenorhabditis elegans protein C42C1.9 (KEO4), mRNA. /FEA=mRNA /GEN=KEO4 /PROD=similar to Caenorhabditis elegans proteinC42C1.9 /DB_XREF=gi:5453705 /UG=Hs.285818 similar to Caenorhabditis elegans protein C42C1.9 /FL=gb:AF064093.1 gb:NM_006459.1	NM_006459		NP_006450
202445_s_at	0.026891	gb:NM_024408.1 /DEF=Homo sapiens Notch (Drosophila) homolog 2 (NOTCH2), mRNA. /FEA=mRNA /GEN=NOTCH2 /PROD=Notch (Drosophila) homolog 2 /DB_XREF=gi:13249343 /UG=Hs.8121 Notch (Drosophila) homolog 2 /FL=gb:AF315356.1 gb:NM_024408.1	NM_024408		NP_077719
202446_s_at	7.58E-04	phospholipid scramblase 1	AI825926	Hs.198282	NP_066928

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202447_at	0.00161	gb:NM_001359.1 /DEF=Homo sapiens 2,4-dienoyl CoA reductase 1, mitochondrial (DEC1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=DEC1 /PROD=2,4-dienoyl CoA reductase 1 precursor /DB_XREF=gi:4503300 /UG=Hs.81548 2,4-dienoyl CoA reductase 1, mitochondrial /FL=gb:U49352.1 gb:NM_001359.1 gb:L26050.1	NM_001359		NP_001350
202457_s_at	7.58E-04	protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha)	AA911231	Hs.272458	NP_000935
202462_s_at	0.00161	gb:NM_014829.1 /DEF=Homo sapiens KIAA0801 gene product (KIAA0801), mRNA. /FEA=mRNA /GEN=KIAA0801 /PROD=KIAA0801 gene product /DB_XREF=gi:7662317 /UG=Hs.17585 KIAA0801 gene product /FL=gb:AB018344.1 gb:NM_014829.1	NM_014829		NP_055644
202464_s_at	2.05E-04	gb:NM_004566.1 /DEF=Homo sapiens 6-phosphofructo-2-kinasefructose-2,6-biphosphatase 3 (PFKFB3), mRNA. /FEA=mRNA /GEN=PFKFB3 /PROD=6-phosphofructo-2-kinasefructose-2,6-biphosphatase 3 /DB_XREF=gi:4758899 /UG=Hs.195471 6-phosphofructo-2-kinasefructose-2,6-biphosphatase 3 /FL=gb:D49817.1 gb:AF109735.1 gb:NM_004566.1	NM_004566		NP_004557
202469_s_at	7.58E-04	cleavage and polyadenylation specific factor 6, 68kDa	AU149367	Hs.64542	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202471_s_at	0.003171	gb:NM_004135.1 /DEF=Homo sapiens isocitrate dehydrogenase 3 (NAD+) gamma (IDH3G), mRNA. /FEA=mRNA /GEN=IDH3G /PROD=isocitrate dehydrogenase 3 (NAD+) gamma /DB_XREF=gi:4758581 /UG=Hs.75253 isocitrate dehydrogenase 3 (NAD+) gamma /FL=gb:BC001902.1 gb:BC000933.2 gb:U40272.1 gb:NM_004135.1	NM_004135		NP_777358
202474_s_at	0.007779	gb:NM_005334.1 /DEF=Homo sapiens host cell factor C1 (VP16-accessory protein) (HCFC1), mRNA. /FEA=mRNA /GEN=HCFC1 /PROD=host cell factor C1 (VP16-accessory protein) /DB_XREF=gi:4885402 /UG=Hs.83634 host cell factor C1 (VP16-accessory protein) /FL=gb:NM_005334.1	NM_005334		NP_005325
202478_at	0.001116	gb:NM_021643.1 /DEF=Homo sapiens GS3955 protein (GS3955), mRNA. /FEA=mRNA /GEN=GS3955 /PROD=GS3955 protein /DB_XREF=gi:11056053 /UG=Hs.155418 GS3955 protein /FL=gb:NM_021643.1 gb:BC002637.1 gb:D87119.1	NM_021643		NP_067675
202479_s_at	0.013088	gb:BC002637.1 /DEF=Homo sapiens, GS3955 protein, clone MGC:3860, mRNA, complete cds. /FEA=mRNA /PROD=GS3955 protein /DB_XREF=gi:12803604 /UG=Hs.155418 GS3955 protein /FL=gb:NM_021643.1 gb:BC002637.1 gb:D87119.1	BC002637		NP_067675

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202480_s_at	6.79E-04	gb:NM_004216.1 /DEF=Homo sapiens death effector domain-containing (DEDD), mRNA. /FEA=mRNA /GEN=DEDD /PROD=death effector domain-containing /DB_XREF=gi:4758143 /UG=Hs.169681 death effector domain-containing /FL=gb:AF083236.1 gb:AF043733.1 gb:AF100341.1 gb:NM_004216.1	NM_004216		NP_127491
202484_s_at	0.001116	gb:AF072242.1 /DEF=Homo sapiens methyl-CpG binding protein MBD2 (MBD2) mRNA, complete cds. /FEA=mRNA /GEN=MBD2 /PROD=methyl-CpG binding protein MBD2 /DB_XREF=gi:3800792 /UG=Hs.25674 methyl-CpG binding domain protein 2 /FL=gb:AF072242.1 gb:NM_003927.2	AF072242		NP_056647
202486_at	0.041795	gb:NM_006796.1 /DEF=Homo sapiens AFG3 (ATPase family gene 3, yeast)-like 2 (AFG3L2), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=AFG3L2 /PROD=AFG3-like 2 /DB_XREF=gi:5802969 /UG=Hs.29385 AFG3 (ATPase family gene 3, yeast)-like 2 /FL=gb:NM_006796.1	NM_006796		NP_006787
202487_s_at	1.26E-04	gb:NM_012412.1 /DEF=Homo sapiens purine-rich element binding protein B (PURB), mRNA. /FEA=mRNA /GEN=PURB /PROD=purine-rich element binding protein B /DB_XREF=gi:6912615 /UG=Hs.301005 purine-rich element binding protein B /FL=gb:BC000098.1 gb:BC004274.1 gb:AF081192.1 gb:NM_012412.1	NM_012412		NP_619541

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
202492_at	0.010205	gb:NM_024085.1 /DEF=Homo sapiens hypothetical protein FLJ22169 (FLJ22169), mRNA. /FEA=mRNA /GEN=FLJ22169 /PROD=hypothetical protein FLJ22169 /DB_XREF=gi:13129081 /UG=Hs.323363 hypothetical protein FLJ22169 /FL=gb:BC001206.1 gb:NM_024085.1	NM_024085		NP_076990
202495_at	1.26E-04	gb:NM_003192.1 /DEF=Homo sapiens tubulin-specific chaperone c (TBCC), mRNA. /FEA=mRNA /GEN=TBCC /PROD=beta-tubulin cofactor C /DB_XREF=gi:4507372 /UG=Hs.75064 tubulin-specific chaperone c /FL=gb:U61234.1 gb:NM_003192.1	NM_003192		NP_003183
202497_x_at	0.041055	Consensus includes gb:AI631159 /FEA=EST /DB_XREF=gi:4682489 /DB_XREF=est:ts93d05.x1 /CLONE=IMAGE:2238825 /UG=Hs.7594 solute carrier family 2 (facilitated glucose transporter), member 3 /FL=gb:M20681.1 gb:NM_006931.1	NM_006931		NP_008862
202499_s_at	2.05E-04	gb:NM_006931.1 /DEF=Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 3 (SLC2A3), mRNA. /FEA=mRNA /GEN=SLC2A3 /PROD=solute carrier family 2 (facilitated glucosetransporter), member 3 /DB_XREF=gi:5902089 /UG=Hs.7594 solute carrier family 2 (facilitated glucose transporter), member 3 /FL=gb:M20681.1 gb:NM_006931.1	NM_006931		NP_008862

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202503_s_at	1.26E-04	gb:NM_014736.1 /DEF=Homo sapiens KIAA0101 gene product (KIAA0101), mRNA. /FEA=mRNA /GEN=KIAA0101 /PROD=KIAA0101 gene product /DB_XREF=gi:7661905 /UG=Hs.81892 KIAA0101 gene product /FL=gb:D14657.1 gb:NM_014736.1	NM_014736		NP_055551
202506_at	5.03E-04	gb:NM_006751.1 /DEF=Homo sapiens sperm specific antigen 2 (SSFA2), mRNA. /FEA=mRNA /GEN=SSFA2 /PROD=sperm specific antigen 2 /DB_XREF=gi:5803178 /UG=Hs.82767 sperm specific antigen 2 /FL=gb:M61199.1 gb:NM_006751.1	NM_006751		NP_006742
202510_s_at	0.041055	gb:NM_006291.1 /DEF=Homo sapiens tumor necrosis factor, alpha-induced protein 2 (TNFAIP2), mRNA. /FEA=mRNA /GEN=TNFAIP2 /PROD=tumor necrosis factor, alpha-induced protein 2 /DB_XREF=gi:5454133 /UG=Hs.101382 tumor necrosis factor, alpha-induced protein 2 /FL=gb:M92357.1 gb:NM_006291.1	NM_006291		NP_006282
202511_s_at	2.05E-04	Consensus includes gb:AK001899.1 /DEF=Homo sapiens cDNA FLJ11037 fis, clone PLACE1004316, highly similar to H.sapiens mRNA for apoptosis specific protein. /FEA=mRNA /DB_XREF=gi:7023451 /UG=Hs.11171 APG5 (autophagy 5, S. cerevisiae)-like /FL=gb:NM_004849.1	AK001899		NP_004840

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202513_s_at	0.004341	gb:NM_006245.1 /DEF=Homo sapiens protein phosphatase 2, regulatory subunit B (B56), delta isoform (PPP2R5D), mRNA. /FEA=mRNA /GEN=PPP2R5D /PROD=protein phosphatase 2, regulatory subunit B(B56), delta isoform /DB_XREF=gi:5453953 /UG=Hs.118244 protein phosphatase 2, regulatory subunit B (B56), delta isoform /FL=gb:BC001095.1 gb:L76702.1 gb:AB000634.1 gb:NM_006245.1	NM_006245		NP_851308
202519_at	0.00228	gb:NM_014938.1 /DEF=Homo sapiens KIAA0867 protein (MONDOA), mRNA. /FEA=mRNA /GEN=MONDOA /PROD=MondoA protein /DB_XREF=gi:7662347 /UG=Hs.52081 KIAA0867 protein /FL=gb:AB020674.1 gb:NM_014938.1	NM_014938		NP_055753
202521_at	0.003171	gb:NM_006565.1 /DEF=Homo sapiens CCCTC-binding factor (zinc finger protein) (CTCF), mRNA. /FEA=mRNA /GEN=CTCF /PROD=CCCTC-binding factor (zinc finger protein) /DB_XREF=gi:5729789 /UG=Hs.57419 CCCTC-binding factor (zinc finger protein) /FL=gb:NM_006565.1 gb:U25435.1	NM_006565		NP_006556
202524_s_at	0.010205	gb:NM_014767.1 /DEF=Homo sapiens KIAA0275 gene product (KIAA0275), mRNA. /FEA=mRNA /GEN=KIAA0275 /PROD=KIAA0275 gene product /DB_XREF=gi:7662035 /UG=Hs.74583 KIAA0275 gene product /FL=gb:D87465.1 gb:NM_014767.1	NM_014767		NP_055582

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202527_s_at	0.00228	gb:NM_005359.1 /DEF=Homo sapiens MAD (mothers against decapentaplegic, Drosophila) homolog 4 (MADH4), mRNA. /FEA=mRNA /GEN=MADH4 /PROD=MAD (mothers against decapentaplegic, Drosophila) homolog 4 /DB_XREF=gi:4885456 /UG=Hs.75862 MAD (mothers against decapentaplegic, Drosophila) homolog 4 /FL=gb:U44378.1 gb:BC002379.1 gb:NM_005359.1	NM_005359		NP_005350
202529_at	5.03E-04	gb:NM_002766.1 /DEF=Homo sapiens phosphoribosyl pyrophosphate synthetase-associated protein 1 (PRPSAP1), mRNA. /FEA=mRNA /GEN=PRPSAP1 /PROD=phosphoribosyl pyrophosphatesynthetase-associated protein 1 /DB_XREF=gi:4506130 /UG=Hs.77498 phosphoribosyl pyrophosphate synthetase-associated protein 1 /FL=gb:D61391.1 gb:NM_002766.1	NM_002766		NP_002757
202530_at	1.26E-04	gb:NM_001315.1 /DEF=Homo sapiens mitogen-activated protein kinase 14 (MAPK14), mRNA. /FEA=mRNA /GEN=MAPK14 /PROD=mitogen-activated protein kinase 14 /DB_XREF=gi:4503068 /UG=Hs.79107 mitogen-activated protein kinase 14 /FL=gb:NM_001315.1 gb:L35263.1	NM_001315		NP_620583
202532_s_at	0.001116	gb:BC000192.1 /DEF=Homo sapiens, dihydrofolate reductase, clone MGC:857, mRNA, complete cds. /FEA=mRNA /PROD=dihydrofolate reductase /DB_XREF=gi:12652874 /UG=Hs.83765 dihydrofolate reductase /FL=gb:BC000192.1 gb:BC003584.1 gb:NM_000791.2	BC000192		NP_000782

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202534_x_at	3.26E-04	gb:NM_000791.2 /DEF=Homo sapiens dihydrofolate reductase (DHFR), mRNA. /FEA=mRNA /GEN=DHFR /PROD=dihydrofolate reductase /DB_XREF=gi:7262376 /UG=Hs.83765 dihydrofolate reductase /FL=gb:BC000192.1 gb:BC003584.1 gb:NM_000791.2	NM_000791		NP_000782
202535_at	2.05E-04	gb:NM_003824.1 /DEF=Homo sapiens Fas (TNFRSF6)-associated via death domain (FADD), mRNA. /FEA=mRNA /GEN=FADD /PROD=Fas (TNFRSF6)-associated via death domain /DB_XREF=gi:4505228 /UG=Hs.86131 Fas (TNFRSF6)-associated via death domain /FL=gb:BC000334.1 gb:NM_003824.1 gb:U24231.1	NM_003824		NP_003815
202536_at	1.26E-04	Consensus includes gb:AK002165.1 /DEF=Homo sapiens cDNA FLJ11303 fis, clone PLACE1009995, highly similar to Homo sapiens mRNA; cDNA DKFZp564O123. /FEA=mRNA /DB_XREF=gi:7023876 /UG=Hs.11449 DKFZP564O123 protein /FL=gb:AF151842.1 gb:AL080122.1 gb:NM_014043.1	AK002165		NP_054762
202538_s_at	1.26E-04	gb:NM_014043.1 /DEF=Homo sapiens DKFZP564O123 protein (DKFZP564O123), mRNA. /FEA=mRNA /GEN=DKFZP564O123 /PROD=DKFZP564O123 protein /DB_XREF=gi:7661633 /UG=Hs.11449 DKFZP564O123 protein /FL=gb:AF151842.1 gb:AL080122.1 gb:NM_014043.1	NM_014043		NP_054762
202539_s_at	1.26E-04	3-hydroxy-3-methylglutaryl-Coenzyme A reductase	AL518627	Hs.11899	NP_000850

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202540_s_at	0.001116	gb:NM_000859.1 /DEF=Homo sapiens 3-hydroxy-3-methylglutaryl-Coenzyme A reductase (HMGCR), mRNA. /FEA=mRNA /GEN=HMGCR /PROD=3-hydroxy-3-methylglutaryl-Coenzyme A reductase /DB_XREF=gi:4557642 /UG=Hs.11899 3-hydroxy-3-methylglutaryl-Coenzyme A reductase /FL=gb:M11058.1 gb:NM_000859.1	NM_000859		NP_000850
202543_s_at	7.58E-04	gb:BC005359.1 /DEF=Homo sapiens, glia maturation factor, beta, clone MGC:12462, mRNA, complete cds. /FEA=mRNA /PROD=glia maturation factor, beta /DB_XREF=gi:13529184 /UG=Hs.151413 glia maturation factor, beta /FL=gb:BC005359.1 gb:M86492.1 gb:AB001106.1 gb:NM_004124.1	BC005359		NP_004115
202544_at	0.005852	gb:NM_004124.1 /DEF=Homo sapiens glia maturation factor, beta (GMFB), mRNA. /FEA=mRNA /GEN=GMFB /PROD=glia maturation factor, beta /DB_XREF=gi:4758441 /UG=Hs.151413 glia maturation factor, beta /FL=gb:BC005359.1 gb:M86492.1 gb:AB001106.1 gb:NM_004124.1	NM_004124		NP_004115
202550_s_at	1.26E-04	gb:NM_004738.1 /DEF=Homo sapiens VAMP (vesicle-associated membrane protein)-associated protein B and C (VAPB), mRNA. /FEA=mRNA /GEN=VAPB /PROD=VAMP (vesicle-associated membraneprotein)-associated protein B and C /DB_XREF=gi:4759301 /UG=Hs.182625 VAMP (vesicle-associated membrane protein)-associated protein B and C /FL=gb:BC001712.1 gb:AF086628.1 gb:NM_004738.1 gb:AF160212.1	NM_004738		NP_004729

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
202553_s_at	0.001116	gb:NM_015484.1 /DEF=Homo sapiens GCIP-interacting protein p29 (P29), mRNA. /FEA=mRNA /GEN=P29 /PROD=GCIP-interacting protein p29 /DB_XREF=gi:7661635 /UG=Hs.20013 GCIP-interacting protein p29 /FL=gb:AF273089.1 gb:AL080166.1 gb:NM_015484.1	NM_015484		NP_056299
202556_s_at	0.02145	gb:NM_006337.1 /DEF=Homo sapiens microspherule protein 1 (MCRS1), mRNA. /FEA=mRNA /GEN=MCRS1 /PROD=microspherule protein 1 /DB_XREF=gi:5453693 /UG=Hs.25313 microspherule protein 1 /FL=gb:AF068007.1 gb:NM_006337.1	NM_006337		NP_006328
202565_s_at	1.26E-04	gb:NM_003174.2 /DEF=Homo sapiens supervillin (SVIL), transcript variant 1, mRNA. /FEA=mRNA /GEN=SVIL /PROD=supervillin, isoform 1 /DB_XREF=gi:11496980 /UG=Hs.154567 supervillin /FL=gb:NM_003174.2 gb:AF051850.1 gb:AF051851.1	NM_003174		NP_068506
202566_s_at	0.016934	gb:AF051851.1 /DEF=Homo sapiens supervillin mRNA, complete cds. /FEA=mRNA /PROD=supervillin /DB_XREF=gi:2961251 /UG=Hs.154567 supervillin /FL=gb:NM_003174.2 gb:AF051850.1 gb:AF051851.1	AF051851		NP_068506
202567_at	0.013223	gb:NM_004175.1 /DEF=Homo sapiens small nuclear ribonucleoprotein D3 polypeptide (18kD) (SNRPD3), mRNA. /FEA=mRNA /GEN=SNRPD3 /PROD=small nuclear ribonucleoprotein D3 polypeptide(18kD) /DB_XREF=gi:4759159 /UG=Hs.1575 small nuclear ribonucleoprotein D3 polypeptide (18kD) /FL=gb:BC000457.1 gb:BC003150.1 gb:NM_004175.1 gb:U15009.1	NM_004175		NP_004166

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202568_s_at	5.03E-04	MAP/microtubule affinity-regulating kinase 3	A1745639	Hs.172766	NP_002367
202569_s_at	0.00161	gb:NM_002376.1 /DEF=Homo sapiens MAPmicrotubule affinity-regulating kinase 3 (MARK3), mRNA. /FEA=mRNA /GEN=MARK3 /PROD=MAPmicrotubule affinity-regulating kinase 3 /DB_XREF=gi:4505102 /UG=Hs.172766 MAPmicrotubule affinity-regulating kinase 3 /FL=gb:M80359.1 gb:NM_002376.1	NM_002376		NP_002367
202582_s_at	1.26E-04	gb:AF306510.1 /DEF=Homo sapiens RANBPM mRNA, complete cds. /FEA=mRNA /PROD=RANBPM /DB_XREF=gi:13194575 /UG=Hs.279886 RAN binding protein 9 /FL=gb:AF306510.1 gb:AB008515.1 gb:NM_005493.1	AF306510		NP_005484
202583_s_at	1.26E-04	gb:NM_005493.1 /DEF=Homo sapiens RAN binding protein 9 (RANBP9), mRNA. /FEA=mRNA /GEN=RANBP9 /PROD=RAN binding protein 9 /DB_XREF=gi:4885570 /UG=Hs.279886 RAN binding protein 9 /FL=gb:AF306510.1 gb:AB008515.1 gb:NM_005493.1	NM_005493		NP_005484
202589_at	1.28E-04	gb:NM_001071.1 /DEF=Homo sapiens thymidylate synthetase (TYMS), mRNA. /FEA=mRNA /GEN=TYMS /PROD=thymidylate synthetase /DB_XREF=gi:4507750 /UG=Hs.82962 thymidylate synthetase /FL=gb:BC002567.1 gb:NM_001071.1	NM_001071		NP_001062

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
202592_at	0.00161	gb:NM_001487.1 /DEF=Homo sapiens GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 1 (GCN5L1), mRNA. /FEA=mRNA /GEN=GCN5L1 /PROD=GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 1 /DB_XREF=gi:4503954 /UG=Hs.94672 GCN5 (general control of amino-acid synthesis, yeast, homolog)-like 1 /FL=gb:D64007.1 gb:NM_001487.1	NM_001487		NP_001478
202593_s_at	0.026891	gb:NM_016641.1 /DEF=Homo sapiens membrane interacting protein of RGS16 (MIR16), mRNA. /FEA=mRNA /GEN=MIR16 /PROD=membrane interacting protein of RGS16 /DB_XREF=gi:7706616 /UG=Hs.107014 membrane interacting protein of RGS16 /FL=gb:AF212862.1 gb:NM_016641.1	NM_016641		NP_057725
202596_at	2.05E-04	gb:BC000436.1 /DEF=Homo sapiens, endosulfine alpha, clone MGC:8394, mRNA, complete cds. /FEA=mRNA /PROD=endosulfine alpha /DB_XREF=gi:12653334 /UG=Hs.111680 endosulfine alpha /FL=gb:BC000436.1 gb:BC004461.1 gb:NM_004436.1 gb:AF157509.1	BC000436		NP_004427
202600_s_at	0.008596	nuclear receptor interacting protein 1	AI824012	Hs.155017	NP_003480
202603_at	3.26E-04	a disintegrin and metalloproteinase domain 10	N51370	Hs.172028	NP_001101

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202604_x_at	0.026891	gb:NM_001110.1 /DEF=Homo sapiens a disintegrin and metalloproteinase domain 10 (ADAM10), mRNA. /FEA=mRNA /GEN=ADAM10 /PROD=a disintegrin and metalloprotease domain 10 /DB_XREF=gi:4557250 /UG=Hs.172028 a disintegrin and metalloproteinase domain 10 /FL=gb:AF009615.1 gb:NM_001110.1	NM_001110		NP_001101
202605_at	0.007779	gb:NM_000181.1 /DEF=Homo sapiens glucuronidase, beta (GUSB), mRNA. /FEA=mRNA /GEN=GUSB /PROD=glucuronidase, beta /DB_XREF=gi:4504222 /UG=Hs.183868 glucuronidase, beta /FL=gb:M15182.1 gb:NM_000181.1	NM_000181		NP_000172
202606_s_at	0.005852	gb:NM_012290.1 /DEF=Homo sapiens tousled-like kinase 1 (TLK1), mRNA. /FEA=mRNA /GEN=TLK1 /PROD=tousled-like kinase 1 /DB_XREF=gi:6912719 /UG=Hs.18895 tousled-like kinase 1 /FL=gb:AB004885.1 gb:NM_012290.1 gb:AF246219.1	NM_012290		NP_036422
202607_at	0.00476	MAD2 mitotic arrest deficient-like 1 (yeast)	AL526632	Hs.79078	NP_001534
202610_s_at	1.26E-04	gb:AF135802.1 /DEF=Homo sapiens thyroid hormone receptor-associated protein complex component TRAP170 mRNA, complete cds. /FEA=mRNA /PROD=thyroid hormone receptor-associated proteincomplex component TRAP170 /DB_XREF=gi:4580325 /UG=Hs.21586 cofactor required for Sp1 transcriptional activation, subunit 2 (150kD) /FL=gb:AF304448.1 gb:AB006651.1 gb:AF104256.1 gb:AF135802.1 gb:NM_004229.1	AF135802		NP_004220
202611_s_at	7.58E-04	cofactor required for Sp1 transcriptional activation, subunit 2, 150kDa	AI971089	Hs.21586	NP_004220

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202615_at	0.00228	guanine nucleotide binding protein (G protein), q polypeptide	BF222895	Hs.380144	
202621_at	1.26E-04	gb:NM_001571.1 /DEF=Homo sapiens interferon regulatory factor 3 (IRF3), mRNA. /FEA=mRNA /GEN=IRF3 /PROD=interferon regulatory factor 3 /DB_XREF=gi:4504724 /UG=Hs.75254 interferon regulatory factor 3 /FL=gb:NM_001571.1	NM_001571		NP_001562
202622_s_at	0.0375	gb:NM_002973.1 /DEF=Homo sapiens spinocerebellar ataxia 2 (olivopontocerebellar ataxia 2, autosomal dominant, ataxin 2) (SCA2), mRNA. /FEA=mRNA /GEN=SCA2 /PROD=ataxin 2 /DB_XREF=gi:4506794 /UG=Hs.76253 spinocerebellar ataxia 2 (olivopontocerebellar ataxia 2, autosomal dominant, ataxin 2) /FL=gb:U70323.1 gb:NM_002973.1	NM_002973		NP_002964
202623_at	0.02145	gb:NM_018453.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM036 (BM036), mRNA. /FEA=mRNA /GEN=BM036 /PROD=uncharacterized bone marrow protein BM036 /DB_XREF=gi:8922092 /UG=Hs.7731 uncharacterized bone marrow protein BM036 /FL=gb:BC001245.1 gb:AF217512.1 gb:NM_018453.1	NM_018453		NP_060923
202625_at	0.026891	v-yes-1 Yamaguchi sarcoma viral related oncogene homolog	AI356412	Hs.80887	NP_002341
202629_at	0.001116	amyloid beta precursor protein (cytoplasmic tail) binding protein 2	AA046411	Hs.84084	NP_006371

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202633_at	0.041055	gb:NM_007027.1 /DEF=Homo sapiens topoisomerase (DNA) II binding protein (TOPBP1), mRNA. /FEA=mRNA /GEN=TOPBP1 /PROD=topoisomerase (DNA) II binding protein /DB_XREF=gi:5902137 /UG=Hs.91417 topoisomerase (DNA) II binding protein /FL=gb:AB019397.1 gb:NM_007027.1	NM_007027		NP_008958
202634_at	0.041055	polymerase (RNA) II (DNA directed) polypeptide K, 7.0kDa	AL558030	Hs.351475	NP_005025
202635_s_at	0.001116	gb:NM_005034.1 /DEF=Homo sapiens polymerase (RNA) II (DNA directed) polypeptide K (7.0kD) (POLR2K), mRNA. /FEA=mRNA /GEN=POLR2K /PROD=polymerase (RNA) II (DNA directed) polypeptide K(7.0kD) /DB_XREF=gi:4826923 /UG=Hs.150675 polymerase (RNA) II (DNA directed) polypeptide K (7.0kD) /FL=gb:BC000806.1 gb:NM_005034.1	NM_005034		NP_005025
202636_at	5.03E-04	gb:NM_005667.1 /DEF=Homo sapiens zinc finger protein homologous to Zfp103 in mouse (ZFP103), mRNA. /FEA=mRNA /GEN=ZFP103 /PROD=zinc finger protein homologous to Zfp103 inmouse /DB_XREF=gi:5031824 /UG=Hs.155968 zinc finger protein homologous to Zfp103 in mouse /FL=gb:D76444.1 gb:NM_005667.1	NM_005667		NP_005658
202637_s_at	0.009179	intercellular adhesion molecule 1 (CD54), human rhinovirus receptor	AI608725	Hs.168383	NP_000192

Gene Identifier	p-value	Description	Gene Accession No.	Unig ne Accession No.	Protein Accession No.
202638_s_at	0.005852	gb:NM_000201.1 /DEF=Homo sapiens intercellular adhesion molecule 1 (CD54), human rhinovirus receptor (ICAM1), mRNA. /FEA=mRNA /GEN=ICAM1 /PROD=intercellular adhesion molecule 1 precursor /DB_XREF=gi:4557877 /UG=Hs.168383 intercellular adhesion molecule 1 (CD54), human rhinovirus receptor /FL=gb:M24283.1 gb:J03132.1 gb:NM_000201.1	NM_000201		NP_000192
202640_s_at	0.009199	gb:NM_003624.1 /DEF=Homo sapiens RAN binding protein 3 (RANBP3), transcript variant RANBP3-a, mRNA. /FEA=mRNA /GEN=RANBP3 /PROD=RAN binding protein 3, isoform RANBP3-a /DB_XREF=gi:4506408 /UG=Hs.176657 RAN binding protein 3 /FL=gb:NM_003624.1	NM_003624		NP_015561
202641_at	7.58E-04	gb:NM_004311.1 /DEF=Homo sapiens ADP-ribosylation factor-like 3 (ARL3), mRNA. /FEA=mRNA /GEN=ARL3 /PROD=ADP-ribosylation factor-like 3 /DB_XREF=gi:4757773 /UG=Hs.182215 ADP-ribosylation factor-like 3 /FL=gb:U07151.1 gb:NM_004311.1	NM_004311		NP_004302
202642_s_at	0.010205	gb:NM_003496.1 /DEF=Homo sapiens transformationtranscription domain-associated protein (TRRAP), mRNA. /FEA=mRNA /GEN=TRRAP /PROD=transformationtranscription domain-associatedprotein /DB_XREF=gi:4507690 /UG=Hs.203952 transformationtranscription domain-associated protein /FL=gb:AF076974.1 gb:NM_003496.1	NM_003496		NP_003487

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202645_s_at	0.00161	gb:NM_000244.1 /DEF=Homo sapiens multiple endocrine neoplasia I (MEN1), mRNA. /FEA=mRNA /GEN=MEN1 /PROD=multiple endocrine neoplasia I /DB_XREF=gi:4557744 /UG=Hs.24297 multiple endocrine neoplasia I /FL=gb:U93236.1 gb:NM_000244.1	NM_000244		NP_570716
202649_x_at	7.58E-04	gb:NM_001022.1 /DEF=Homo sapiens ribosomal protein S19 (RPS19), mRNA. /FEA=mRNA /GEN=RPS19 /PROD=ribosomal protein S19 /DB_XREF=gi:4506694 /UG=Hs.298262 ribosomal protein S19 /FL=gb:BC000023.1 gb:M81757.1 gb:NM_001022.1	NM_001022		NP_001013
202650_s_at	0.00161	gb:NM_014738.1 /DEF=Homo sapiens KIAA0195 gene product (KIAA0195), mRNA. /FEA=mRNA /GEN=KIAA0195 /PROD=KIAA0195 gene product /DB_XREF=gi:7661985 /UG=Hs.301132 KIAA0195 gene product /FL=gb:D83779.1 gb:NM_014738.1	NM_014738		NP_055553
202651_at	1.26E-04	gb:NM_014873.1 /DEF=Homo sapiens KIAA0205 gene product (KIAA0205), mRNA. /FEA=mRNA /GEN=KIAA0205 /PROD=KIAA0205 gene product /DB_XREF=gi:7661995 /UG=Hs.3610 KIAA0205 gene product /FL=gb:D86960.1 gb:NM_014873.1	NM_014873		NP_055688

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202653_s_at	0.010205	Consensus includes gb:BC003404.1 /DEF=Homo sapiens, hypothetical protein DKFZp586F1122 similar to axotrophin, clone IMAGE:3449089, mRNA, partial cds. /FEA=mRNA /PROD=hypothetical protein DKFZp586F1122 similar to axotrophin /DB_XREF=gi:13097302 /UG=Hs.5306 hypothetical protein DKFZp586F1122 similar to axotrophin /FL=gb:NM_022826.1	BC003404		NP_073737
202654_x_at	7.58E-04	gb:NM_022826.1 /DEF=Homo sapiens hypothetical protein DKFZp586F1122 similar to axotrophin (DKFZP586F1122), mRNA. /FEA=mRNA /GEN=DKFZP586F1122 /PROD=hypothetical protein DKFZp586F1122 similar to axotrophin /DB_XREF=gi:12383065 /UG=Hs.5306 hypothetical protein DKFZp586F1122 similar to axotrophin /FL=gb:NM_022826.1	NM_022826		NP_073737
202656_s_at	7.58E-04	transcriptional regulator interacting with the PHS-bromodomain 2	BG107456	Hs.77293	NP_055570
202657_s_at	1.26E-04	gb:NM_014755.1 /DEF=Homo sapiens KIAA0127 gene product (KIAA0127), mRNA. /FEA=mRNA /GEN=KIAA0127 /PROD=KIAA0127 gene product /DB_XREF=gi:7661925 /UG=Hs.77293 KIAA0127 gene product /FL=gb:D50917.1 gb:NM_014755.1	NM_014755		NP_055570

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202658_at	0.013223	gb:NM_003846.1 /DEF=Homo sapiens peroxisomal biogenesis factor 11B (PEX11B), mRNA. /FEA=mRNA /GEN=PEX11B /PROD=peroxisomal biogenesis factor 11B /DB_XREF=gi:4505718 /UG=Hs.83023 peroxisomal biogenesis factor 11B /FL=gb:AF093670.1 gb:AB018080.1 gb:NM_003846.1	NM_003846		NP_003837
202660_at	0.001116	inositol 1,4,5-triphosphate receptor, type 2	AA834576	Hs.238272	
202662_s_at	0.005852	gb:NM_002223.1 /DEF=Homo sapiens inositol 1,4,5-triphosphate receptor, type 2 (ITPR2), mRNA. /FEA=mRNA /GEN=ITPR2 /PROD=inositol 1,4,5-triphosphate receptor, type 2 /DB_XREF=gi:4504792 /UG=Hs.238272 inositol 1,4,5-triphosphate receptor, type 2 /FL=gb:D26350.1 gb:NM_002223.1	NM_002223		
202663_at	0.001116	Wiskott-Aldrich syndrome protein interacting protein	AI005043	Hs.379203	
202665_s_at	0.005852	gb:NM_003387.2 /DEF=Homo sapiens Wiskott-Aldrich syndrome protein interacting protein (WASPIP), mRNA. /FEA=mRNA /GEN=WASPIP /PROD=WASP-interacting protein /DB_XREF=gi:8400739 /UG=Hs.24143 Wiskott-Aldrich syndrome protein interacting protein /FL=gb:NM_003387.2	NM_003387		NP_003378
202670_at	1.26E-04	mitogen-activated protein kinase kinase 1	AI571419	Hs.3446	NP_002746
202671_s_at	3.26E-04	gb:NM_003681.1 /DEF=Homo sapiens pyridoxal (pyridoxine, vitamin B6) kinase (PDXK), mRNA. /FEA=mRNA /GEN=PDXK /PROD=pyridoxal kinase /DB_XREF=gi:4505700 /UG=Hs.38041 pyridoxal (pyridoxine, vitamin B6) kinase /FL=gb:BC000123.1 gb:U89606.1 gb:NM_003681.1	NM_003681		NP_003672

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202675_at	1.26E-04	gb:NM_003000.1 /DEF=Homo sapiens succinate dehydrogenase complex, subunit B, iron sulfur (lp) (SDHB), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=SDHB /PROD=succinate dehydrogenase complex, subunit B, iron sulfur (lp) /DB_XREF=gi:9257241 /UG=Hs.64 succinate dehydrogenase complex, subunit B, iron sulfur (lp) /FL=gb:U17248.1 gb:NM_003000.1	NM_003000		NP_002991
202678_at	1.26E-04	gb:NM_004492.1 /DEF=Homo sapiens general transcription factor IIA, 2 (12kD subunit) (GTF2A2), mRNA. /FEA=mRNA /GEN=GTF2A2 /PROD=general transcription factor IIA, 2 (12kD subunit) /DB_XREF=gi:4758485 /UG=Hs.76362 general transcription factor IIA, 2 (12kD subunit) /FL=gb:BC000287.1 gb:BC001919.1 gb:U21242.1 gb:NM_004492.1 gb:U14193.1	NM_004492		NP_004483
202680_at	1.36E-04	gb:NM_002095.1 /DEF=Homo sapiens general transcription factor IIE, polypeptide 2 (beta subunit, 34kD) (GTF2E2), mRNA. /FEA=mRNA /GEN=GTF2E2 /PROD=general transcription factor IIE, polypeptide 2(beta subunit, 34kD) /DB_XREF=gi:4504194 /UG=Hs.77100 general transcription factor IIE, polypeptide 2 (beta subunit, 34kD) /FL=gb:NM_002095.1	NM_002095		NP_002086

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202687_s_at	7.58E-04	gb:U57059.1 /DEF=Homo sapiens Apo-2 ligand mRNA, complete cds. /FEA=mRNA /PROD=Apo-2 ligand /DB_XREF=gi:1336207 /UG=Hs.83429 tumor necrosis factor (ligand) superfamily, member 10 /FL=gb:U37518.1 gb:U57059.1 gb:NM_003810.1	U57059		NP_003801
202688_at	0.016934	gb:NM_003810.1 /DEF=Homo sapiens tumor necrosis factor (ligand) superfamily, member 10 (TNFSF10), mRNA. /FEA=mRNA /GEN=TNFSF10 /PROD=tumor necrosis factor (ligand) superfamily, member 10 /DB_XREF=gi:4507592 /UG=Hs.83429 tumor necrosis factor (ligand) superfamily, member 10 /FL=gb:U37518.1 gb:U57059.1 gb:NM_003810.1	NM_003810		NP_003801
202690_s_at	0.005852	gb:BC001721.1 /DEF=Homo sapiens, small nuclear ribonucleoprotein D1 polypeptide (16kD), clone MGC:2222, mRNA, complete cds. /FEA=mRNA /PROD=small nuclear ribonucleoprotein D1 polypeptide(16kD) /DB_XREF=gi:12804598 /UG=Hs.86948 small nuclear ribonucleoprotein D1 polypeptide (16kD) /FL=gb:BC001721.1 gb:J03798.1 gb:NM_006938.1	BC001721		NP_008869
202695_s_at	0.013223	gb:NM_004760.1 /DEF=Homo sapiens serinethreonine kinase 17a (apoptosis-inducing) (STK17A), mRNA. /FEA=mRNA /GEN=STK17A /PROD=serinethreonine kinase 17a(apoptosis-inducing) /DB_XREF=gi:4758191 /UG=Hs.9075 serinethreonine kinase 17a (apoptosis-inducing) /FL=gb:AB011420.1 gb:NM_004760.1	NM_004760		NP_004751

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202696_at	1.26E-04	gb:NM_005109.1 /DEF=Homo sapiens oxidative-stress responsive 1 (OSR1), mRNA. /FEA=mRNA /GEN=OSR1 /PROD=oxidative-stress responsive 1 /DB_XREF=gi:4826877 /UG=Hs.95220 oxidative-stress responsive 1 /FL=gb:AB017642.1 gb:NM_005109.1 gb:AB029024.1	NM_005109		NP_005100
202697_at	1.26E-04	gb:NM_007006.1 /DEF=Homo sapiens cleavage and polyadenylation specific factor 5, 25 kD subunit (CPSF5), mRNA. /FEA=mRNA /GEN=CPSF5 /PROD=cleavage and polyadenylation specific factor 5,25 kD subunit /DB_XREF=gi:5901925 /UG=Hs.9605 cleavage and polyadenylation specific factor 5, 25 kD subunit /FL=gb:BC001403.1 gb:NM_007006.1	NM_007006		NP_008937
202700_s_at	0.004541	gb:NM_014698.1 /DEF=Homo sapiens KIAA0792 gene product (KIAA0792), mRNA. /FEA=mRNA /GEN=KIAA0792 /PROD=KIAA0792 gene product /DB_XREF=gi:7662307 /UG=Hs.119387 KIAA0792 gene product /FL=gb:AB018335.1 gb:NM_014698.1	NM_014698		NP_055513
202708_s_at	0.004341	gb:NM_003528.1 /DEF=Homo sapiens H2B histone family, member Q (H2BFQ), mRNA. /FEA=mRNA /GEN=H2BFQ /PROD=H2B histone family, member Q /DB_XREF=gi:4504276 /UG=Hs.2178 H2B histone family, member Q /FL=gb:NM_003528.1	NM_003528		NP_003519

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202719_s_at	0.0174	gb:BC001451.1 /DEF=Homo sapiens, testin, clone MGC:1146, mRNA, complete cds. /FEA=mRNA /PROD=testin /DB_XREF=gi:12655188 /UG=Hs.165986 testin /FL=gb:AF245356.1 gb:AF245357.1 gb:BC001451.1 gb:NM_015641.1	BC001451		NP_690042
202720_at	0.02145	gb:NM_015641.1 /DEF=Homo sapiens testin (DKFZP586B2022), mRNA. /FEA=mRNA /GEN=DKFZP586B2022 /PROD=testin /DB_XREF=gi:7661665 /UG=Hs.165986 testin /FL=gb:AF245356.1 gb:AF245357.1 gb:BC001451.1 gb:NM_015641.1	NM_015641		NP_690042
202725_at	2.05E-04	gb:NM_000937.1 /DEF=Homo sapiens polymerase (RNA) II (DNA directed) polypeptide A (220kD) (POLR2A), mRNA. /FEA=mRNA /GEN=POLR2A /PROD=polymerase (RNA) II (DNA directed) polypeptide A(220kD) /DB_XREF=gi:4505938 /UG=Hs.171880 polymerase (RNA) II (DNA directed) polypeptide A (220kD) /FL=gb:NM_000937.1	NM_000937		NP_000928
202727_s_at	3.26E-04	gb:NM_000416.1 /DEF=Homo sapiens interferon gamma receptor 1 (IFNGR1), mRNA. /FEA=mRNA /GEN=IFNGR1 /PROD=interferon gamma receptor 1 /DB_XREF=gi:4557879 /UG=Hs.180866 interferon gamma receptor 1 /FL=gb:BC005333.1 gb:J03143.1 gb:NM_000416.1	NM_000416		NP_000407

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202729_s_at	0.046687	gb:NM_000627.1 /DEF=Homo sapiens latent transforming growth factor beta binding protein 1 (LTBP1), mRNA. /FEA=mRNA /GEN=LTBP1 /PROD=latent transforming growth factor beta bindingprotein 1 precursor /DB_XREF=gi:4557730 /UG=Hs.241257 latent transforming growth factor beta binding protein 1 /FL=gb:M34057.1 gb:NM_000627.1	NM_000627		NP_000618
202730_s_at	0.02145	gb:NM_014456.1 /DEF=Homo sapiens programmed cell death 4 (PDCD4), mRNA. /FEA=mRNA /GEN=PDCD4 /PROD=programmed cell death 4 /DB_XREF=gi:7657448 /UG=Hs.296251 programmed cell death 4 /FL=gb:U96628.1 gb:NM_014456.1	NM_014456		NP_663314
202731_at	0.004341	gb:NM_014456.1 /DEF=Homo sapiens programmed cell death 4 (PDCD4), mRNA. /FEA=mRNA /GEN=PDCD4 /PROD=programmed cell death 4 /DB_XREF=gi:7657448 /UG=Hs.296251 programmed cell death 4 /FL=gb:U96628.1 gb:NM_014456.1	NM_014456		NP_663314
202738_s_at	0.005852	phosphorylase kinase, beta	BG149218	Hs.78060	NP_000284
202739_s_at	0.016934	gb:NM_000293.1 /DEF=Homo sapiens phosphorylase kinase, beta (PHKB), mRNA. /FEA=mRNA /GEN=PHKB /PROD=phosphorylase kinase, beta /DB_XREF=gi:4505782 /UG=Hs.78060 phosphorylase kinase, beta /FL=gb:NM_000293.1	NM_000293		NP_000284
202741_at	0.026891	protein kinase, cAMP-dependent, catalytic, beta	AA130247	Hs.87773	NP_002722

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202742_s_at	0.00228	gb:NM_002731.1 /DEF=Homo sapiens protein kinase, cAMP-dependent, catalytic, beta (PRKACB), mRNA. /FEA=mRNA /GEN=PRKACB /PROD=protein kinase, cAMP-dependent, catalytic, beta /DB_XREF=gi:4506056 /UG=Hs.87773 protein kinase, cAMP-dependent, catalytic, beta /FL=gb:M34181.1 gb:NM_002731.1	NM_002731		NP_002722
202745_at	0.016934	gb:NM_005154.1 /DEF=Homo sapiens ubiquitin specific protease 8 (USP8), mRNA. /FEA=mRNA /GEN=USP8 /PROD=ubiquitin specific protease 8 /DB_XREF=gi:4827053 /UG=Hs.152818 ubiquitin specific protease 8 /FL=gb:D29956.1 gb:NM_005154.1	NM_005154		NP_005145
202749_at	0.016934	gb:NM_004627.1 /DEF=Homo sapiens tryptophan rich basic protein (WRB), mRNA. /FEA=mRNA /GEN=WRB /PROD=tryptophan rich basic protein /DB_XREF=gi:4759325 /UG=Hs.198308 tryptophan rich basic protein /FL=gb:NM_004627.1	NM_004627		NP_004618
202750_s_at	0.016686	gb:AL080147.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434B194 (from clone DKFZp434B194); complete cds. /FEA=mRNA /GEN=DKFZp434B194 /PROD=hypothetical protein /DB_XREF=gi:5262598 /UG=Hs.20225 tuftelin-interacting protein /FL=gb:AL080147.1 gb:NM_012143.1	AL080147		NP_036275

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202753_at	0.004341	gb:NM_014814.1 /DEF=Homo sapiens KIAA0107 gene product (KIAA0107), mRNA. /FEA=mRNA /GEN=KIAA0107 /PROD=KIAA0107 gene product /DB_XREF=gi:7661913 /UG=Hs.23488 KIAA0107 gene product /FL=gb:BC000630.1 gb:BC000904.2 gb:D14663.1 gb:AF215935.1 gb:NM_014814.1	NM_014814		NP_055629
202758_s_at	0.003928	gb:NM_003721.1 /DEF=Homo sapiens regulatory factor X-associated ankyrin-containing protein (RFXANK), mRNA. /FEA=mRNA /GEN=RFXANK /PROD=regulatory factor X-associatedankyrin-containing protein /DB_XREF=gi:4506498 /UG=Hs.296776 regulatory factor X-associated ankyrin-containing protein /FL=gb:AF094760.1 gb:AF105427.1 gb:NM_003721.1 gb:AF077196.1	NM_003721		NP_604389
202761_s_at	0.041055	gb:NM_015180.1 /DEF=Homo sapiens synaptic nuclei expressed gene 2; KIAA1011 protein (KIAA1011), mRNA. /FEA=mRNA /GEN=KIAA1011 /PROD=KIAA1011 protein /DB_XREF=gi:11056019 /UG=Hs.57749 synaptic nuclei expressed gene 2; KIAA1011 protein /FL=gb:NM_015180.1 gb:AL080133.1	NM_015180		
202763_at	0.026891	gb:NM_004346.1 /DEF=Homo sapiens caspase 3, apoptosis-related cysteine protease (CASP3), mRNA. /FEA=mRNA /GEN=CASP3 /PROD=caspase 3, apoptosis-related cysteine protease /DB_XREF=gi:4757911 /UG=Hs.74552 caspase 3, apoptosis-related cysteine protease /FL=gb:NM_004346.1 gb:U13737.1 gb:U13738.1 gb:U26943.1	NM_004346		NP_116786

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202764_at	0.001054	gb:NM_003156.1 /DEF=Homo sapiens stromal interaction molecule 1 (STIM1), mRNA. /FEA=mRNA /GEN=STIM1 /PROD=stromal interaction molecule 1 /DB_XREF=gi:4507268 /UG=Hs.74597 stromal interaction molecule 1 /FL=gb:U52426.1 gb:NM_003156.1	NM_003156		NP_003147
202767_at	7.58E-04	gb:NM_001610.1 /DEF=Homo sapiens acid phosphatase 2, lysosomal (ACP2), mRNA. /FEA=mRNA /GEN=ACP2 /PROD=lysosomal acid phosphatase 2 precursor /DB_XREF=gi:4557009 /UG=Hs.75589 acid phosphatase 2, lysosomal /FL=gb:BC003160.1 gb:NM_001610.1	NM_001610		NP_001601
202770_s_at	5.03E-04	gb:NM_004354.1 /DEF=Homo sapiens cyclin G2 (CCNG2), mRNA. /FEA=mRNA /GEN=CCNG2 /PROD=cyclin G2 /DB_XREF=gi:4757935 /UG=Hs.79069 cyclin G2 /FL=gb:U47414.1 gb:NM_004354.1	NM_004354		NP_004345
202771_at	0.029453	gb:NM_014745.1 /DEF=Homo sapiens KIAA0233 gene product (KIAA0233), mRNA. /FEA=mRNA /GEN=KIAA0233 /PROD=KIAA0233 gene product /DB_XREF=gi:7662013 /UG=Hs.79077 KIAA0233 gene product /FL=gb:D87071.1 gb:NM_014745.1	NM_014745		NP_055560

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202772_at	0.018123	gb:NM_000191.1 /DEF=Homo sapiens 3-hydroxymethyl-3-methylglutaryl-Coenzyme A lyase (hydroxymethylglutaricaciduria) (HMGCL), mRNA. /FEA=mRNA /GEN=HMGCL /PROD=3-hydroxymethyl-3-methylglutaryl-Coenzyme A lyase (hydroxymethylglutaricaciduria) /DB_XREF=gi:4504426 /UG=Hs.831 3-hydroxymethyl-3-methylglutaryl-Coenzyme A lyase (hydroxymethylglutaricaciduria) /FL=gb:L07033.1 gb:NM_000191.1	NM_000191		NP_000182
202774_s_at	0.013088	splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot homolog, Drosophila)	AI023864	Hs.84229	NP_689421
202777_at	1.26E-04	gb:NM_007373.1 /DEF=Homo sapiens suppressor of clear, C. elegans, homolog of (SHOC2), mRNA. /FEA=mRNA /GEN=SHOC2 /PROD=suppressor of clear, C. elegans, homolog of /DB_XREF=gi:6677944 /UG=Hs.104315 soc-2 (suppressor of clear, C.elegans) homolog /FL=gb:AF068920.1 gb:AF054828.1 gb:AB020669.1 gb:NM_007373.1	NM_007373		NP_031399
202778_s_at	0.026891	gb:NM_003453.1 /DEF=Homo sapiens zinc finger protein 198 (ZNF198), mRNA. /FEA=mRNA /GEN=ZNF198 /PROD=zinc finger protein 198 /DB_XREF=gi:4508010 /UG=Hs.109526 zinc finger protein 198 /FL=gb:AF035374.1 gb:AF060181.1 gb:NM_003453.1	NM_003453		NP_003444

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202782_s_at	0.007052	gb:NM_016532.1 /DEF=Homo sapiens SKIP for skeletal muscle and kidney enriched inositol phosphatase (LOC51763), mRNA. /FEA=mRNA /GEN=LOC51763 /PROD=SKIP for skeletal muscle and kidney enriched inositol phosphatase /DB_XREF=gi:7706564 /UG=Hs.178347 SKIP for skeletal muscle and kidney enriched inositol phosphatase /FL=gb:AB036829.1 gb:NM_016532.1	NM_016532		NP_570122
202788_at	2.05E-04	gb:NM_004635.1 /DEF=Homo sapiens mitogen-activated protein kinase-activated protein kinase 3 (MAPKAPK3), mRNA. /FEA=mRNA /GEN=MAPKAPK3 /PROD=mitogen-activated protein kinase-activated protein kinase 3 /DB_XREF=gi:4758699 /UG=Hs.227789 mitogen-activated protein kinase-activated protein kinase 3 /FL=gb:U09578.1 gb:U43784.1 gb:BC001662.1 gb:NM_004635.1	NM_004635		NP_004626
202794_at	0.00228	gb:NM_002194.2 /DEF=Homo sapiens inositol polyphosphate-1-phosphatase (INPP1), mRNA. /FEA=mRNA /GEN=INPP1 /PROD=inositol polyphosphate-1-phosphatase /DB_XREF=gi:4755138 /UG=Hs.32309 inositol polyphosphate-1-phosphatase /FL=gb:L08488.1 gb:NM_002194.2	NM_002194		NP_002185
202795_x_at	0.041055	gb:NM_007032.1 /DEF=Homo sapiens putative nuclear protein (HRIHFB2122), mRNA. /FEA=mRNA /GEN=HRIHFB2122 /PROD=putative nuclear protein /DB_XREF=gi:10334853 /UG=Hs.40342 putative nuclear protein /FL=gb:NM_007032.1	NM_007032		NP_619538

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202798_at	0.005852	gb:NM_006323.1 /DEF=Homo sapiens SEC24 (S. cerevisiae) related gene family, member B (SEC24B), mRNA. /FEA=mRNA /GEN=SEC24B /PROD=SEC24 (S. cerevisiae) related gene family, member B /DB_XREF=gi:5454045 /UG=Hs.7239 SEC24 (S. cerevisiae) related gene family, member B /FL=gb:NM_006323.1	NM_006323		NP_006314
2028_s_at	0.002864	E2F transcription factor 1	NM_005225	Hs.96055	NP_005216
202801_at	0.041055	gb:NM_002730.1 /DEF=Homo sapiens protein kinase, cAMP-dependent, catalytic, alpha (PRKACA), mRNA. /FEA=mRNA /GEN=PRKACA /PROD=protein kinase, cAMP-dependent, catalytic, alpha /DB_XREF=gi:4506054 /UG=Hs.77271 protein kinase, cAMP-dependent, catalytic, alpha /FL=gb:NM_002730.1	NM_002730		NP_002721
202807_s_at	0.001054	gb:NM_005488.1 /DEF=Homo sapiens target of myb1 (chicken) homolog (TOM1), mRNA. /FEA=mRNA /GEN=TOM1 /PROD=target of myb1 (chicken) homolog /DB_XREF=gi:4885636 /UG=Hs.9482 target of myb1 (chicken) homolog /FL=gb:NM_005488.1	NM_005488		NP_005479
202810_at	0.016934	gb:NM_004147.1 /DEF=Homo sapiens developmentally regulated GTP-binding protein 1 (DRG1), mRNA. /FEA=mRNA /GEN=DRG1 /PROD=developmentally regulated GTP-binding protein 1 /DB_XREF=gi:4758795 /UG=Hs.115242 developmentally regulated GTP-binding protein 1 /FL=gb:AF078103.1 gb:NM_004147.1	NM_004147		NP_004138

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202811_at	5.03E-04	gb:NM_006463.1 /DEF=Homo sapiens associated molecule with the SH3 domain of STAM (AMSH), mRNA. /FEA=mRNA /GEN=AMSH /PROD=associated molecule with the SH3 domain of STAM /DB_XREF=gi:5453544 /UG=Hs.12479 associated molecule with the SH3 domain of STAM /FL=gb:U73522.1 gb:NM_006463.1	NM_006463		NP_006454
202814_s_at	0.016934	HMBA-inducible	AW193511	Hs.15299	NP_006451
202823_at	0.036254	transcription elongation factor B (SIII), polypeptide 1 (15kDa, elongin C)	N89607	Hs.184693	NP_005639
202824_s_at	3.26E-04	gb:NM_005648.1 /DEF=Homo sapiens transcription elongation factor B (SIII), polypeptide 1 (15kD, elongin C) (TCEB1), mRNA. /FEA=mRNA /GEN=TCEB1 /PROD=elongin C /DB_XREF=gi:5032160 /UG=Hs.184693 transcription elongation factor B (SIII), polypeptide 1 (15kD, elongin C) /FL=gb:NM_005648.1 gb:L34587.1	NM_005648		NP_005639
202829_s_at	1.26E-04	gb:NM_005638.1 /DEF=Homo sapiens synaptobrevin-like 1 (SYBL1), mRNA. /FEA=mRNA /GEN=SYBL1 /PROD=synaptobrevin-like 1 /DB_XREF=gi:5032136 /UG=Hs.24167 synaptobrevin-like 1 /FL=gb:NM_005638.1	NM_005638		NP_005629

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202833_s_at	1.26E-04	gb:NM_000295.1 /DEF=Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1 (SERPINA1), mRNA. /FEA=mRNA /GEN=SERPINA1 /PROD=serine (or cysteine) proteinase inhibitor, cladeA (alpha 1 antiproteinase, antitrypsin), member 1 /DB_XREF=gi:4505792 /UG=Hs.297681 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1 /FL=gb:AF130068.1 gb:M11465.1 gb:K01396.1 gb:NM_000295.1	NM_000295		NP_000286
202836_s_at	7.58E-04	gb:NM_006701.1 /DEF=Homo sapiens similar to S. pombe dim1+ (DIM1), mRNA. /FEA=mRNA /GEN=DIM1 /PROD=similar to S. pombe dim1+ /DB_XREF=gi:5729801 /UG=Hs.5074 similar to S. pombe dim1+ /FL=gb:BC001046.1 gb:AF023611.1 gb:NM_006701.1 gb:AF146373.1	NM_006701		NP_006692
202845_s_at	3.26E-04	gb:NM_006788.1 /DEF=Homo sapiens ralA binding protein 1 (RALBP1), mRNA. /FEA=mRNA /GEN=RALBP1 /PROD=ralA binding protein 1 /DB_XREF=gi:5803144 /UG=Hs.75447 ralA binding protein 1 /FL=gb:NM_006788.1 gb:L42542.1	NM_006788		NP_006779
202848_s_at	0.016934	G protein-coupled receptor kinase 6	BG423052	Hs.76297	NP_002073

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202852_s_at	5.07E-04	gb:NM_024666.1 /DEF=Homo sapiens hypothetical protein FLJ11506 (FLJ11506), mRNA. /FEA=mRNA /GEN=FLJ11506 /PROD=hypothetical protein FLJ11506 /DB_XREF=gi:13375923 /UG=Hs.77703 hypothetical protein FLJ11506 /FL=gb:AL136715.1 gb:NM_024666.1	NM_024666		NP_078942
202854_at	1.26E-04	gb:NM_000194.1 /DEF=Homo sapiens hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan syndrome) (HPRT1), mRNA. /FEA=mRNA /GEN=HPRT1 /PROD=hypoxanthine phosphoribosyltransferase 1 /DB_XREF=gi:4504482 /UG=Hs.82314 hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan syndrome) /FL=gb:BC000578.1 gb:M31642.1 gb:NM_000194.1	NM_000194		NP_000185
202859_x_at	0.00228	gb:NM_000584.1 /DEF=Homo sapiens interleukin 8 (IL8), mRNA. /FEA=mRNA /GEN=IL8 /PROD=interleukin 8 /DB_XREF=gi:10834977 /UG=Hs.624 interleukin 8 /FL=gb:NM_000584.1 gb:M17017.1 gb:M26383.1	NM_000584		NP_000575
202872_at	2.05E-04	Consensus includes gb:AW024925 /FEA=EST /DB_XREF=gi:5878455 /DB_XREF=est:wu92g09.x1 /CLONE=IMAGE:2527552 /UG=Hs.86905 ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 42kD /FL=gb:NM_001695.1	NM_001695		NP_001686

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202873_at	0.03002	Consensus includes gb:BF034973 /FEA=EST /DB_XREF=gi:10742685 /DB_XREF=est:601459039F1 /CLONE=IMAGE:3862832 /UG=Hs.86905 ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD /FL=gb:NM_001695.1	NM_001695		NP_001686
202874_s_at	0.00161	gb:NM_001695.1 /DEF=Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD (ATP6C), mRNA. /FEA=mRNA /GEN=ATP6C /PROD=ATPase, H+ transporting, lysosomal (vacuolarproton pump) 42kD /DB_XREF=gi:4502314 /UG=Hs.86905 ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD /FL=gb:NM_001695.1	NM_001695		NP_001686
202878_s_at	2.05E-04	gb:NM_012072.2 /DEF=Homo sapiens complement component C1q receptor (C1QR), mRNA. /FEA=mRNA /GEN=C1QR /PROD=complement component C1q receptor /DB_XREF=gi:11496985 /UG=Hs.97199 complement component C1q receptor /FL=gb:NM_012072.2 gb:U94333.1	NM_012072		NP_036204
202880_s_at	0.003171	gb:NM_004762.1 /DEF=Homo sapiens pleckstrin homology, Sec7 and coiledcoil domains 1(cytohesin 1) (PSCD1), transcript variant 1, mRNA. /FEA=mRNA /GEN=PSCD1 /PROD=cytohesin 1, isoform 1 /DB_XREF=gi:4758963 /UG=Hs.1050 pleckstrin homology, Sec7 and coiledcoil domains 1(cytohesin 1) /FL=gb:M85169.1 gb:NM_004762.1 gb:NM_017456.1	NM_004762		NP_059430

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202882_x_at	0.007779	gb:NM_016167.2 /DEF=Homo sapiens retinoic acid repressible protein (RARG-1), mRNA. /FEA=mRNA /GEN=RARG-1 /PROD=retinoic acid repressible protein /DB_XREF=gi:13569842 /UG=Hs.106346 retinoic acid repressible protein /FL=gb:AF172066.2 gb:NM_016167.2	NM_016167		NP_057251
202888_s_at	0.007779	gb:NM_001150.1 /DEF=Homo sapiens alanyl (membrane) aminopeptidase (aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD13, p150) (ANPEP), mRNA. /FEA=mRNA /GEN=ANPEP /PROD=membrane alanine aminopeptidase precursor /DB_XREF=gi:4502094 /UG=Hs.1239 alanyl (membrane) aminopeptidase (aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD13, p150) /FL=gb:M22324.1 gb:NM_001150.1	NM_001150		NP_001141
202891_at	1.26E-04	gb:NM_005600.1 /DEF=Homo sapiens nitrilase 1 (NIT1), mRNA. /FEA=mRNA /GEN=NIT1 /PROD=nitrilase 1 /DB_XREF=gi:5031946 /UG=Hs.146406 nitrilase 1 /FL=gb:AF069987.1 gb:NM_005600.1	NM_005600		NP_005591
202897_at	1.26E-04	gb:AB023430.1 /DEF=Homo sapiens Bit mRNA, complete cds. /FEA=mRNA /GEN=Bit /DB_XREF=gi:6518912 /UG=Hs.156114 protein tyrosine phosphatase, non-receptor type substrate 1 /FL=gb:D86043.1 gb:NM_004648.1 gb:AB023430.1	AB023430		NP_542970

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202899_s_at	0.041055	gb:NM_003017.1 /DEF=Homo sapiens splicing factor, arginineserine-rich 3 (SFRS3), mRNA. /FEA=mRNA /GEN=SFRS3 /PROD=splicing factor, arginineserine-rich 3 /DB_XREF=gi:4506900 /UG=Hs.167460 splicing factor, arginineserine-rich 3 /FL=gb:L10838.1 gb:NM_003017.1	NM_003017		NP_003008
202907_s_at	0.00228	gb:NM_002485.2 /DEF=Homo sapiens Nijmegen breakage syndrome 1 (nibrin) (NBS1), mRNA. /FEA=mRNA /GEN=NBS1 /PROD=nibrin /DB_XREF=gi:6996019 /UG=Hs.25812 Nijmegen breakage syndrome 1 (nibrin) /FL=gb:AF058696.1 gb:AF051334.1 gb:NM_002485.2	NM_002485		NP_002476
202912_at	0.033381	gb:NM_001124.1 /DEF=Homo sapiens adrenomedullin (ADM), mRNA. /FEA=mRNA /GEN=ADM /PROD=adrenomedullin /DB_XREF=gi:4501944 /UG=Hs.394 adrenomedullin /FL=gb:NM_001124.1 gb:D14874.1	NM_001124		NP_001115
202916_s_at	0.016402	gb:NM_014864.1 /DEF=Homo sapiens KIAA0475 gene product (KIAA0475), mRNA. /FEA=mRNA /GEN=KIAA0475 /PROD=KIAA0475 gene product /DB_XREF=gi:7662149 /UG=Hs.5737 KIAA0475 gene product /FL=gb:AB007944.1 gb:NM_014864.1	NM_014864		NP_055679
202918_s_at	0.005852	gb:AF151853.1 /DEF=Homo sapiens CGI-95 protein mRNA, complete cds. /FEA=mRNA /PROD=CGI-95 protein /DB_XREF=gi:4929658 /UG=Hs.107942 DKFZP564M112 protein /FL=gb:AB015441.1 gb:BC005237.1 gb:AF151853.1 gb:AL080070.1 gb:NM_015387.1	AF151853		NP_056202

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202925_s_at	7.58E-04	gb:NM_002657.2 /DEF=Homo sapiens pleiomorphic adenoma gene-like 2 (PLAGL2), mRNA. /FEA=mRNA /GEN=PLAGL2 /PROD=pleiomorphic adenoma gene-like 2 /DB_XREF=gi:6031195 /UG=Hs.154104 pleiomorphic adenoma gene-like 2 /FL=gb:AF006005.1 gb:NM_002657.2	NM_002657		NP_002648
202934_at	1.26E-04	hexokinase 2	AI761561	Hs.198427	NP_000180
202939_at	3.26E-04	gb:NM_005857.1 /DEF=Homo sapiens zinc metalloproteinase, STE24 (yeast, homolog) (ZMPSTE24), mRNA. /FEA=mRNA /GEN=ZMPSTE24 /PROD=zinc metalloproteinase, STE24 (yeast, homolog) /DB_XREF=gi:5032128 /UG=Hs.25846 zinc metalloproteinase, STE24 (yeast, homolog) /FL=gb:AB016068.1 gb:AF064867.1 gb:NM_005857.1	NM_005857		NP_005848
202941_at	7.58E-04	gb:NM_021074.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) flavoprotein 2 (24kD) (NDUFV2), mRNA. /FEA=mRNA /GEN=NDUFV2 /PROD=NADH dehydrogenase (ubiquinone) flavoprotein 2(24kD) /DB_XREF=gi:10835024 /UG=Hs.51299 NADH dehydrogenase (ubiquinone) flavoprotein 2 (24kD) /FL=gb:NM_021074.1 gb:BC001632.1 gb:M22538.1	NM_021074		NP_066552
202944_at	0.010205	gb:NM_000262.1 /DEF=Homo sapiens N-acetylgalactosaminidase, alpha- (NAGA), mRNA. /FEA=mRNA /GEN=NAGA /PROD=alpha-N-acetylgalactosaminidase precursor /DB_XREF=gi:4557780 /UG=Hs.75372 N-acetylgalactosaminidase, alpha- /FL=gb:BC000095.1 gb:M62783.1 gb:M38083.1 gb:NM_000262.1	NM_000262		NP_000253

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202946_s_at	1.92E-04	gb:NM_014962.1 /DEF=Homo sapiens KIAA0952 protein (KIAA0952), mRNA. /FEA=mRNA /GEN=KIAA0952 /PROD=KIAA0952 protein /DB_XREF=gi:7662401 /UG=Hs.7935 KIAA0952 protein /FL=gb:AB023169.1 gb:NM_014962.1	NM_014962		NP_852108
202947_s_at	2.76E-04	gb:NM_002101.2 /DEF=Homo sapiens glycophorin C (Gerbich blood group) (GYPC), transcript variant 1, mRNA. /FEA=mRNA /GEN=GYPC /PROD=glycophorin C, isoform 1 /DB_XREF=gi:8051606 /UG=Hs.81994 glycophorin C (Gerbich blood group) /FL=gb:M36284.1 gb:M11802.1 gb:NM_002101.2	NM_002101		NP_058131
202948_at	1.28E-04	gb:NM_000877.1 /DEF=Homo sapiens interleukin 1 receptor, type I (IL1R1), mRNA. /FEA=mRNA /GEN=IL1R1 /PROD=interleukin 1 receptor, type I /DB_XREF=gi:4504658 /UG=Hs.82112 interleukin 1 receptor, type I /FL=gb:M27492.1 gb:NM_000877.1	NM_000877		NP_000868
202951_at	0.041055	serine/threonine kinase 38	BE048506	Hs.8724	NP_009202
202954_at	0.023856	gb:NM_007019.1 /DEF=Homo sapiens ubiquitin carrier protein E2-C (UBCH10), mRNA. /FEA=mRNA /GEN=UBCH10 /PROD=ubiquitin carrier protein E2-C /DB_XREF=gi:5902145 /UG=Hs.93002 ubiquitin carrier protein E2-C /FL=gb:U73379.1 gb:NM_007019.1	NM_007019		NP_008950

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202955_s_at	0.027243	gb:AF084520.1 /DEF=Homo sapiens brefeldin A-inhibited guanine nucleotide-exchange protein 1 mRNA, complete cds. /FEA=mRNA /PROD=brefeldin A-inhibited guaninenucleotide-exchange protein 1 /DB_XREF=gi:5052120 /UG=Hs.94631 brefeldin A-inhibited guanine nucleotide-exchange protein 1 /FL=gb:AF084520.1 gb:AF111162.1 gb:NM_006421.2	AF084520		NP_006412
202956_at	5.03E-04	gb:NM_006421.2 /DEF=Homo sapiens brefeldin A-inhibited guanine nucleotide-exchange protein 1 (BIG1), mRNA. /FEA=mRNA /GEN=BIG1 /PROD=brefeldin A-inhibited guaninenucleotide-exchange protein 1 /DB_XREF=gi:6715588 /UG=Hs.94631 brefeldin A-inhibited guanine nucleotide-exchange protein 1 /FL=gb:AF084520.1 gb:AF111162.1 gb:NM_006421.2	NM_006421		NP_006412
202958_at	0.046435	gb:NM_002833.1 /DEF=Homo sapiens protein tyrosine phosphatase, non-receptor type 9 (PTPN9), mRNA. /FEA=mRNA /GEN=PTPN9 /PROD=protein tyrosine phosphatase, non-receptor type9 /DB_XREF=gi:4506300 /UG=Hs.147663 protein tyrosine phosphatase, non-receptor type 9 /FL=gb:M83738.1 gb:NM_002833.1	NM_002833		NP_002824

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202960_s_at	0.007779	gb:NM_000255.1 /DEF=Homo sapiens methylmalonyl Coenzyme A mutase (MUT), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=MUT /PROD=methylmalonyl Coenzyme A mutase precursor /DB_XREF=gi:4557766 /UG=Hs.155212 methylmalonyl Coenzyme A mutase /FL=gb:M65131.1 gb:NM_000255.1	NM_000255		NP_000246
202961_s_at	7.58E-04	gb:NM_004889.1 /DEF=Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit f, isoform 2 (ATP5J2), mRNA. /FEA=mRNA /GEN=ATP5J2 /PROD=ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit f, isoform 2 /DB_XREF=gi:4757811 /UG=Hs.155751 ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit f, isoform 2 /FL=gb:BC003678.1 gb:AF047436.1 gb:NM_004889.1	NM_004889		NP_004880
202968_s_at	0.008036	Consensus includes gb:Y09216.1 /DEF=H.sapiens mRNA for protein kinase, Dyrk2. /FEA=mRNA /GEN=Dyrk2 /DB_XREF=gi:1666065 /UG=Hs.173135 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 /FL=gb:NM_006482.1	Y09216		NP_006473
202969_at	0.001116	Consensus includes gb:AI216690 /FEA=EST /DB_XREF=gi:3785731 /DB_XREF=est:qg66h07.x1 /CLONE=IMAGE:1840189 /UG=Hs.173135 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 /FL=gb:NM_006482.1	Y09216		NP_006473

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202971_s_at	0.018693	gb:NM_006482.1 /DEF=Homo sapiens dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 (DYRK2), transcript variant 2, mRNA. /FEA=mRNA /GEN=DYRK2 /PROD=dual-specificity tyrosine-(Y)-phosphorylationregulated kinase 2 isoform 2 /DB_XREF=gi:5922003 /UG=Hs.173135 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 /FL=gb:NM_006482.1	NM_006482		NP_006473
202972_s_at	0.018206	KIAA0914 gene product	AW450403	Hs.177664	NP_055698
202974_at	1.26E-04	gb:NM_002436.2 /DEF=Homo sapiens membrane protein, palmitoylated 1 (55kD) (MPP1), mRNA. /FEA=mRNA /GEN=MPP1 /PROD=palmitoylated membrane protein 1 /DB_XREF=gi:6006024 /UG=Hs.1861 membrane protein, palmitoylated 1 (55kD) /FL=gb:BC002392.1 gb:M64925.1 gb:NM_002436.2	NM_002436		NP_002427
202981_x_at	0.033381	gb:NM_003031.1 /DEF=Homo sapiens seven in absentia (Drosophila) homolog 1 (SIAH1), mRNA. /FEA=mRNA /GEN=SIAH1 /PROD=seven in absentia (Drosophila) homolog 1 /DB_XREF=gi:4506946 /UG=Hs.295923 seven in absentia (Drosophila) homolog 1 /FL=gb:U76247.1 gb:U63295.1 gb:NM_003031.1	NM_003031		NP_003022

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
202990_at	1.26E-04	gb:NM_002863.1 /DEF=Homo sapiens phosphorylase, glycogen; liver (Hers disease, glycogen storage disease type VI) (PYGL), mRNA. /FEA=mRNA /GEN=PYGL /PROD=phosphorylase, glycogen; liver (Hers disease, glycogen storage disease type VI) /DB_XREF=gi:4506352 /UG=Hs.771 phosphorylase, glycogen; liver (Hers disease, glycogen storage disease type VI) /FL=gb:M14636.1 gb:AF066858.1 gb:AF046785.1 gb:NM_002863.1	NM_002863		NP_002854
202995_s_at	6.42E-04	gb:NM_006486.1 /DEF=Homo sapiens fibulin 1 (FBLN1), transcript variant D, mRNA. /FEA=mRNA /GEN=FBLN1 /PROD=fibulin 1 isoform D /DB_XREF=gi:5922008 /UG=Hs.79732 fibulin 1 /FL=gb:U01244.1 gb:NM_006486.1	NM_006486		NP_006478
203003_at	0.035171	MADS box transcription enhancer factor 2, polypeptide D (myocyte enhancer factor 2D)	AL530331	Hs.111243	NP_005911
203005_at	1.26E-04	gb:NM_002342.1 /DEF=Homo sapiens lymphotoxin beta receptor (TNFR superfamily, member 3 (LTBR), mRNA. /FEA=mRNA /GEN=LTBR /PROD=lymphotoxin B receptor /DB_XREF=gi:4505038 /UG=Hs.1116 lymphotoxin beta receptor (TNFR superfamily, member 3 /FL=gb:L04270.1 gb:NM_002342.1	NM_002342		NP_002333

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203008_x_at	0.005852	gb:NM_005783.1 /DEF=Homo sapiens ATP binding protein associated with cell differentiation (APACD), mRNA. /FEA=mRNA /GEN=APACD /PROD=ATP binding protein associated with celldifferentiation /DB_XREF=gi:5031582 /UG=Hs.153884 ATP binding protein associated with cell differentiation /FL=gb:AB006679.1 gb:NM_005783.1	NM_005783		NP_005774
203011_at	0.001116	gb:NM_005536.2 /DEF=Homo sapiens inositol(myo)-1(or 4)-monophosphatase 1 (IMPA1), mRNA. /FEA=mRNA /GEN=IMPA1 /PROD=inositol(myo)-1(or 4)-monophosphatase 1 /DB_XREF=gi:8393607 /UG=Hs.171776 inositol(myo)-1(or 4)-monophosphatase 1 /FL=gb:AF042729.2 gb:NM_005536.2	NM_005536		NP_005527
203012_x_at	0.013223	gb:NM_000984.1 /DEF=Homo sapiens ribosomal protein L23a (RPL23A), mRNA. /FEA=mRNA /GEN=RPL23A /PROD=ribosomal protein L23a /DB_XREF=gi:4506614 /UG=Hs.184776 ribosomal protein L23a /FL=gb:U37230.1 gb:NM_000984.1	NM_000984		NP_000975
203013_at	0.021448	gb:NM_007265.1 /DEF=Homo sapiens suppressor of S. cerevisiae gcr2 (HSGT1), mRNA. /FEA=mRNA /GEN=HSGT1 /PROD=suppressor of S. cerevisiae gcr2 /DB_XREF=gi:6005783 /UG=Hs.19673 suppressor of S. cerevisiae gcr2 /FL=gb:BC000721.1 gb:D88208.1 gb:NM_007265.1	NM_007265		NP_009196
203017_s_at	0.027239	KIAA0923 protein	AW136988	Hs.22587	NP_054740

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203021_at	2.05E-04	gb:NM_003064.1 /DEF=Homo sapiens secretory leukocyte protease inhibitor (antileukoproteinase) (SLPI), mRNA. /FEA=mRNA /GEN=SLPI /PROD=secretory leukocyte protease inhibitor(antileukoproteinase) /DB_XREF=gi:4507064 /UG=Hs.251754 secretory leukocyte protease inhibitor (antileukoproteinase) /FL=gb:NM_003066.1 gb:AF114471.1 gb:NM_003064.1	NM_003064		NP_003055
203024_s_at	0.033381	gb:NM_020199.1 /DEF=Homo sapiens HTGN29 protein (HTGN29), mRNA. /FEA=mRNA /GEN=HTGN29 /PROD=HTGN29 protein /DB_XREF=gi:9910277 /UG=Hs.283437 HTGN29 protein /FL=gb:AF226055.1 gb:NM_020199.1	NM_020199		NP_064584
203025_at	0.009199	gb:NM_003491.1 /DEF=Homo sapiens N-acetyltransferase, homolog of S. cerevisiae ARD1 (ARD1), mRNA. /FEA=mRNA /GEN=ARD1 /PROD=N-acetyltransferase, homolog of S. cerevisiaeARD1 /DB_XREF=gi:10835056 /UG=Hs.333034 N-acetyltransferase, homolog of S. cerevisiae ARD1 /FL=gb:NM_003491.1 gb:BC000308.1	NM_003491		NP_003482

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203039_s_at	0.001116	gb:NM_005006.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 1 (75kD) (NADH-coenzyme Q reductase) (NDUFS1), mRNA. /FEA=mRNA /GEN=NDUFS1 /PROD=NADH dehydrogenase (ubiquinone) Fe-S protein 1(75kD) (NADH-coenzyme Q reductase) /DB_XREF=gi:4826855 /UG=Hs.8248 NADH dehydrogenase (ubiquinone) Fe-S protein 1 (75kD) (NADH-coenzyme Q reductase) /FL=gb:NM_005006.1	NM_005006		NP_004997
203041_s_at	1.26E-04	gb:J04183.1 /DEF=Homo sapiens lysosomal membrane glycoprotein-2 (LAMP2), complete cds. /FEA=mRNA /GEN=LAMP2 /PROD=lysosomal membrane glycoprotein-2 /DB_XREF=gi:186929 /UG=Hs.8262 lysosomal-associated membrane protein 2 /FL=gb:J04183.1 gb:NM_002294.1	J04183		NP_054701
203042_at	2.05E-04	gb:NM_002294.1 /DEF=Homo sapiens lysosomal-associated membrane protein 2 (LAMP2), transcript variant LAMP2A, mRNA. /FEA=mRNA /GEN=LAMP2 /PROD=lysosomal-associated membrane protein 2precursor /DB_XREF=gi:4504956 /UG=Hs.8262 lysosomal-associated membrane protein 2 /FL=gb:J04183.1 gb:NM_002294.1	NM_002294		NP_054701
203044_at	3.26E-04	gb:NM_014918.1 /DEF=Homo sapiens KIAA0990 protein (KIAA0990), mRNA. /FEA=mRNA /GEN=KIAA0990 /PROD=KIAA0990 protein /DB_XREF=gi:7662433 /UG=Hs.110488 KIAA0990 protein /FL=gb:AB023207.1 gb:NM_014918.1	NM_014918		NP_055733

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203045_at	0.022795	gb:NM_004148.1 /DEF=Homo sapiens ninjurin 1 (NINJ1), mRNA. /FEA=mRNA /GEN=NINJ1 /PROD=ninjurin 1 /DB_XREF=gi:4758809 /UG=Hs.11342 ninjurin 1 /FL=gb:BC004440.1 gb:U72661.1 gb:U91512.1 gb:NM_004148.1	NM_004148		NP_004139
203060_s_at	0.021045	gb:AF074331.1 /DEF=Homo sapiens PAPS synthetase-2 (PAPSS2) mRNA, complete cds. /FEA=mRNA /GEN=PAPSS2 /PROD=PAPS synthetase-2 /DB_XREF=gi:5052074 /UG=Hs.274230 3-phosphoadenosine 5-phosphosulfate synthase 2 /FL=gb:AF150754.2 gb:AF313907.1 gb:AF091242.1 gb:NM_004670.1 gb:AF074331.1 gb:AF173365.1	AF074331		NP_004661
203064_s_at	0.011176	gb:NM_004514.1 /DEF=Homo sapiens interleukin enhancer binding factor 1 (ILF1), mRNA. /FEA=mRNA /GEN=ILF1 /PROD=interleukin enhancer binding factor 1 /DB_XREF=gi:4758599 /UG=Hs.296281 interleukin enhancer binding factor 1 /FL=gb:U58196.1 gb:NM_004514.1	NM_004514		NP_004505
203074_at	0.02145	gb:NM_001630.1 /DEF=Homo sapiens annexin A8 (ANXA8), mRNA. /FEA=mRNA /GEN=ANXA8 /PROD=annexin VIII /DB_XREF=gi:4502112 /UG=Hs.87268 annexin A8 /FL=gb:BC004376.1 gb:M81844.1 gb:NM_001630.1	NM_001630		NP_001621
203075_at	3.26E-04	MAD, mothers against decapentaplegic homolog 2 (Drosophila)	AW151617	Hs.82483	NP_005892

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203076_s_at	1.26E-04	gb:U65019.1 /DEF=Human MAD-related protein 2 (MADR2) mRNA, complete cds. /FEA=mRNA /GEN=MADR2 /PROD=MAD-related protein 2 /DB_XREF=gi:1575529 /UG=Hs.82483 MAD (mothers against decapentaplegic, Drosophila) homolog 2 /FL=gb:U59911.1 gb:U68018.1 gb:U65019.1 gb:AF027964.1 gb:NM_005901.1	U65019		NP_005892
203077_s_at	0.007779	gb:NM_005901.1 /DEF=Homo sapiens MAD (mothers against decapentaplegic, Drosophila) homolog 2 (MADH2), mRNA. /FEA=mRNA /GEN=MADH2 /PROD=MAD (mothers against decapentaplegic, Drosophila) homolog 2 /DB_XREF=gi:5174510 /UG=Hs.82483 MAD (mothers against decapentaplegic, Drosophila) homolog 2 /FL=gb:U59911.1 gb:U68018.1 gb:U65019.1 gb:AF027964.1 gb:NM_005901.1	NM_005901		NP_005892
203082_at	0.007779	gb:NM_014753.1 /DEF=Homo sapiens KIAA0187 gene product (KIAA0187), mRNA. /FEA=mRNA /GEN=KIAA0187 /PROD=KIAA0187 gene product /DB_XREF=gi:7661979 /UG=Hs.10848 KIAA0187 gene product /FL=gb:D80009.1 gb:NM_014753.1	NM_014753		NP_055568
203090_at	3.26E-04	gb:NM_006923.1 /DEF=Homo sapiens stromal cell-derived factor 2 (SDF2), mRNA. /FEA=mRNA /GEN=SDF2 /PROD=stromal cell-derived factor 2 /DB_XREF=gi:5902073 /UG=Hs.118684 stromal cell-derived factor 2 /FL=gb:BC000500.1 gb:BC001406.1 gb:D50645.1 gb:NM_006923.1	NM_006923		NP_008854

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203094_at	0.013223	gb:NM_014628.1 /DEF=Homo sapiens gene predicted from cDNA with a complete coding sequence (KIAA0110), mRNA. /FEA=mRNA /GEN=KIAA0110 /PROD=gene predicted from cDNA with a complete codingsequence /DB_XREF=gi:7661917 /UG=Hs.124 gene predicted from cDNA with a complete coding sequence /FL=gb:BC002904.1 gb:D14811.1 gb:NM_014628.1	NM_014628		NP_055443
203096_s_at	0.037567	PDZ domain containing guanine nucleotide exchange factor(GEF)1	BF439282	Hs.154545	
203097_s_at	0.003171	gb:NM_014247.1 /DEF=Homo sapiens PDZ domain containing guanine nucleotide exchange factor(GEF)1 (PDZ-GEF1), mRNA. /FEA=mRNA /GEN=PDZ-GEF1 /PROD=PDZ domain containing guanine nucleotideexchange factor(GEF)1 /DB_XREF=gi:7657260 /UG=Hs.154545 PDZ domain containing guanine nucleotide exchange factor(GEF)1 /FL=gb:AB002311.1 gb:NM_014247.1	NM_014247		NP_055062
203100_s_at	0.006603	gb:NM_004824.1 /DEF=Homo sapiens chromodomain protein, Y chromosome-like (CDYL), mRNA. /FEA=mRNA /GEN=CDYL /PROD=chromodomain protein, Y chromosome-like /DB_XREF=gi:4757969 /UG=Hs.16081 chromodomain protein, Y chromosome-like /FL=gb:AF081258.1 gb:AF081259.1 gb:NM_004824.1	NM_004824		NP_736608

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203102_s_at	0.007779	gb:NM_002408.2 /DEF=Homo sapiens mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase (MGAT2), mRNA. /FEA=mRNA /GEN=MGAT2 /PROD=alpha-1,6-mannosyl-glycoproteinbeta-1,2-N-acetylglucosaminyltransferase /DB_XREF=gi:6031183 /UG=Hs.172195 mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase /FL=gb:NM_002408.2	NM_002408		NP_002399
203107_x_at	7.58E-04	gb:NM_002952.1 /DEF=Homo sapiens ribosomal protein S2 (RPS2), mRNA. /FEA=mRNA /GEN=RPS2 /PROD=ribosomal protein S2 /DB_XREF=gi:4506718 /UG=Hs.182426 ribosomal protein S2 /FL=gb:BC001795.1 gb:NM_002952.1	NM_002952		NP_002943
203110_at	0.00161	gb:U43522.1 /DEF=Human cell adhesion kinase beta (CAKbeta) mRNA, complete cds. /FEA=mRNA /GEN=CAKbeta /PROD=cell adhesion kinase beta /DB_XREF=gi:1165218 /UG=Hs.20313 protein tyrosine kinase 2 beta /FL=gb:L49207.1 gb:U43522.1 gb:NM_004103.1 gb:U33284.1	U43522		NP_775268
203111_s_at	0.031424	gb:NM_004103.1 /DEF=Homo sapiens protein tyrosine kinase 2 beta (PTK2B), mRNA. /FEA=mRNA /GEN=PTK2B /PROD=protein tyrosine kinase 2 beta /DB_XREF=gi:4758975 /UG=Hs.20313 protein tyrosine kinase 2 beta /FL=gb:L49207.1 gb:U43522.1 gb:NM_004103.1 gb:U33284.1	NM_004103		NP_775268

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203113_s_at	1.26E-04	gb:NM_001960.1 /DEF=Homo sapiens eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) (EEF1D), mRNA. /FEA=mRNA /GEN=EEF1D /PROD=eukaryotic translation elongation factor 1 delta(guanine nucleotide exchange protein) /DB_XREF=gi:4503478 /UG=Hs.223241 eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) /FL=gb:NM_001960.1	NM_001960		NP_115754
203116_s_at	1.28E-04	gb:NM_000140.1 /DEF=Homo sapiens ferrochelatase (protoporphyria) (FECH), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=FECH /PROD=ferrochelatase precursor /DB_XREF=gi:4557592 /UG=Hs.26 ferrochelatase (protoporphyria) /FL=gb:NM_000140.1	NM_000140		NP_000131
203120_at	1.26E-04	gb:NM_005426.1 /DEF=Homo sapiens tumor protein p53-binding protein, 2 (TP53BP2), mRNA. /FEA=mRNA /GEN=TP53BP2 /PROD=tumor protein p53-binding protein, 2 /DB_XREF=gi:4885642 /UG=Hs.44585 tumor protein p53-binding protein, 2 /FL=gb:U58334.1 gb:NM_005426.1	NM_005426		NP_005417
203123_s_at	0.007779	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2	AU154469	Hs.57435	NP_000608

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203126_at	3.26E-04	gb:NM_014214.1 /DEF=Homo sapiens inositol(myo)-1(or 4)-monophosphatase 2 (IMPA2), mRNA. /FEA=mRNA /GEN=IMPA2 /PROD=inositol(myo)-1(or 4)-monophosphatase 2 /DB_XREF=gi:7657235 /UG=Hs.5753 inositol(myo)-1(or 4)-monophosphatase 2 /FL=gb:AF200432.1 gb:NM_014214.1 gb:AF014398.2	NM_014214		NP_055029
203127_s_at	2.05E-04	gb:BC005123.1 /DEF=Homo sapiens, serine palmitoyltransferase, long chain base subunit 2, clone MGC:10362, mRNA, complete cds. /FEA=mRNA /PROD=serine palmitoyltransferase, long chain basesubunit 2 /DB_XREF=gi:13477298 /UG=Hs.59403 serine palmitoyltransferase, long chain base subunit 2 /FL=gb:BC005123.1 gb:AB011098.1 gb:NM_004863.1	BC005123		NP_004854
203128_at	0.01436	gb:NM_004863.1 /DEF=Homo sapiens serine palmitoyltransferase, long chain base subunit 2 (SPTLC2), mRNA. /FEA=mRNA /GEN=SPTLC2 /PROD=serine palmitoyltransferase, long chain basesubunit 2 /DB_XREF=gi:4758667 /UG=Hs.59403 serine palmitoyltransferase, long chain base subunit 2 /FL=gb:BC005123.1 gb:AB011098.1 gb:NM_004863.1	NM_004863		NP_004854

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203132_at	0.013223	gb:NM_000321.1 /DEF=Homo sapiens retinoblastoma 1 (including osteosarcoma) (RB1), mRNA. /FEA=mRNA /GEN=RB1 /PROD=retinoblastoma 1 (including osteosarcoma) /DB_XREF=gi:4506434 /UG=Hs.75770 retinoblastoma 1 (including osteosarcoma) /FL=gb:M33647.1 gb:M15400.1 gb:M28419.1 gb:NM_000321.1	NM_000321		NP_000312
203133_at	3.26E-04	gb:NM_006808.1 /DEF=Homo sapiens protein translocation complex beta (SEC61B), mRNA. /FEA=mRNA /GEN=SEC61B /PROD=protein translocation complex beta /DB_XREF=gi:5803164 /UG=Hs.77028 protein translocation complex beta /FL=gb:BC001734.1 gb:L25085.1 gb:NM_006808.1	NM_006808		NP_006799
203134_at	0.001747	gb:NM_007166.1 /DEF=Homo sapiens Clathrin assembly lymphoid-myeloid leukemia gene (CLTH), mRNA. /FEA=mRNA /GEN=CLTH /PROD=Clathrin assembly lymphoid-myeloid leukemia gene /DB_XREF=gi:6005732 /UG=Hs.7885 phosphatidylinositol binding clathrin assembly protein /FL=gb:U45976.1 gb:NM_007166.1	NM_007166		NP_009097
203137_at	0.003171	gb:NM_004906.1 /DEF=Homo sapiens Wilms tumour 1-associating protein (KIAA0105), mRNA. /FEA=mRNA /GEN=KIAA0105 /PROD=Wilms tumour 1-associating protein /DB_XREF=gi:4758635 /UG=Hs.119 Wilms tumour 1-associating protein /FL=gb:AF277190.1 gb:D14661.1 gb:NM_004906.1	NM_004906		NP_690597

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203138_at	5.03E-04	gb:NM_003642.1 /DEF=Homo sapiens histone acetyltransferase 1 (HAT1), mRNA. /FEA=mRNA /GEN=HAT1 /PROD=histone acetyltransferase 1 /DB_XREF=gi:4504340 /UG=Hs.13340 histone acetyltransferase 1 /FL=gb:AF030424.1 gb:NM_003642.1	NM_003642		NP_003633
203140_at	0.004341	gb:NM_001706.1 /DEF=Homo sapiens B-cell CLLlymphoma 6 (zinc finger protein 51) (BCL6), mRNA. /FEA=mRNA /GEN=BCL6 /PROD=B-cell CLLlymphoma 6 (zinc finger protein 51) /DB_XREF=gi:4502382 /UG=Hs.155024 B-cell CLLlymphoma 6 (zinc finger protein 51) /FL=gb:U00115.1 gb:NM_001706.1	NM_001706		NP_620309
203141_s_at	0.003395	adaptor-related protein complex 3, beta 1 subunit	AW058575	Hs.155172	NP_003655
203142_s_at	1.26E-04	gb:NM_003664.1 /DEF=Homo sapiens adaptor-related protein complex 3, beta 1 subunit (AP3B1), mRNA. /FEA=mRNA /GEN=AP3B1 /PROD=adaptor-related protein complex 3, beta 1 subunit /DB_XREF=gi:4501974 /UG=Hs.155172 adaptor-related protein complex 3, beta 1 subunit /FL=gb:U91931.1 gb:U81504.1 gb:NM_003664.1	NM_003664		NP_003655
203143_s_at	0.026891	KIAA0040 gene product	T79953	Hs.158282	NP_055471
203144_s_at	0.021244	gb:NM_014656.1 /DEF=Homo sapiens KIAA0040 gene product (KIAA0040), mRNA. /FEA=mRNA /GEN=KIAA0040 /PROD=KIAA0040 gene product /DB_XREF=gi:7657258 /UG=Hs.158282 KIAA0040 gene product /FL=gb:D25539.1 gb:NM_014656.1	NM_014656		NP_055471

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203145_at	0.021312	gb:NM_006461.1 /DEF=Homo sapiens mitotic spindle coiled-coil related protein (DEEPEST), mRNA. /FEA=mRNA /GEN=DEEPEST /PROD=mitotic spindle coiled-coil related protein /DB_XREF=gi:5453631 /UG=Hs.16244 mitotic spindle coiled-coil related protein /FL=gb:BC000322.1 gb:AF063308.1 gb:NM_006461.1	NM_006461		NP_006452
203146_s_at	0.016275	gb:NM_001470.1 /DEF=Homo sapiens gamma-aminobutyric acid (GABA) B receptor, 1 (GABBR1), transcript variant 1, mRNA. /FEA=mRNA /GEN=GABBR1 /PROD=gamma-aminobutyric acid (GABA) B receptor 1, isoform a precursor /DB_XREF=gi:10835014 /UG=Hs.167017 gamma-aminobutyric acid (GABA) B receptor, 1 /FL=gb:NM_001470.1 gb:AF301005.1 gb:AF099148.1	NM_001470		NP_068705
203147_s_at	0.026891	tripartite motif-containing 14	BE962483	Hs.179703	NP_150090
203155_at	0.026891	gb:NM_012432.1 /DEF=Homo sapiens SET domain, bifurcated 1 (SETDB1), mRNA. /FEA=mRNA /GEN=SETDB1 /PROD=SET domain, bifurcated 1 /DB_XREF=gi:6912651 /UG=Hs.20991 SET domain, bifurcated 1 /FL=gb:D31891.1 gb:NM_012432.1	NM_012432		NP_036564
203158_s_at	0.046329	gb:AF097493.1 /DEF=Homo sapiens glutaminase kidney isoform mRNA, complete cds. /FEA=mRNA /PROD=glutaminase kidney isoform /DB_XREF=gi:6002672 /UG=Hs.239189 glutaminase /FL=gb:AF327434.1 gb:AB020645.1 gb:AF097493.1 gb:AF223943.1 gb:NM_014905.1	AF097493		NP_055720

Gene Identifier	p-valu	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203162_s_at	0.02145	gb:NM_005886.1 /DEF=Homo sapiens katanin p80 (WD40-containing) subunit B 1 (KATNB1), mRNA. /FEA=mRNA /GEN=KATNB1 /PROD=katanin p80 subunit B 1 /DB_XREF=gi:5031816 /UG=Hs.275675 katanin p80 (WD40-containing) subunit B 1 /FL=gb:BC001353.1 gb:AF052432.1 gb:NM_005886.1	NM_005886		NP_005877
203165_s_at	0.006698	gb:NM_004733.2 /DEF=Homo sapiens acetyl-Coenzyme A transporter (ACATN), mRNA. /FEA=mRNA /GEN=ACATN /PROD=acetyl-Coenzyme A transporter /DB_XREF=gi:6042194 /UG=Hs.285176 acetyl-Coenzyme A transporter /FL=gb:D88152.1 gb:NM_004733.2	NM_004733		NP_004724
203166_at	0.007779	gb:NM_006324.1 /DEF=Homo sapiens craniofacial development protein 1 (CFDP1), mRNA. /FEA=mRNA /GEN=CFDP1 /PROD=craniofacial development protein 1 /DB_XREF=gi:5453566 /UG=Hs.296460 craniofacial development protein 1 /FL=gb:BC000991.2 gb:AB009285.1 gb:NM_006324.1	NM_006324		NP_006315
203167_at	1.26E-04	gb:NM_003255.2 /DEF=Homo sapiens tissue inhibitor of metalloproteinase 2 (TIMP2), mRNA. /FEA=mRNA /GEN=TIMP2 /PROD=tissue inhibitor of metalloproteinase 2precursor /DB_XREF=gi:9257247 /UG=Hs.325495 tissue inhibitor of metalloproteinase 2 /FL=gb:M32304.1 gb:J05593.1 gb:NM_003255.2	NM_003255		NP_003246

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203184_at	3.26E-04	gb:NM_001999.2 /DEF=Homo sapiens fibrillin 2 (congenital contractural arachnodactyly) (FBN2), mRNA. /FEA=mRNA /GEN=FBN2 /PROD=fibrillin 2 /DB_XREF=gi:4755135 /UG=Hs.79432 fibrillin 2 (congenital contractural arachnodactyly) /FL=gb:U03272.1 gb:NM_001999.2	NM_001999		NP_001990
203185_at	0.00228	gb:NM_014737.1 /DEF=Homo sapiens Ras association (RalGDSAF-6) domain family 2 (RASSF2), mRNA. /FEA=mRNA /GEN=RASSF2 /PROD=Ras association (RalGDSAF-6) domain family 2 /DB_XREF=gi:7661963 /UG=Hs.80905 Ras association (RalGDSAF-6) domain family 2 /FL=gb:D79990.1 gb:NM_014737.1	NM_014737		NP_739580
203186_s_at	0.00228	gb:NM_002961.2 /DEF=Homo sapiens S100 calcium-binding protein A4 (calcium protein, calvasculin, metastasin, murine placental homolog) (S100A4), transcript variant 1, mRNA. /FEA=mRNA /GEN=S100A4 /PROD=S100 calcium-binding protein A4 /DB_XREF=gi:9845514 /UG=Hs.81256 S100 calcium-binding protein A4 (calcium protein, calvasculin, metastasin, murine placental homolog) /FL=gb:NM_002961.2 gb:NM_019554.1	NM_002961		NP_062427
203195_s_at	0.033381	gb:NM_005387.2 /DEF=Homo sapiens nucleoporin 98kD (NUP98), mRNA. /FEA=mRNA /GEN=NUP98 /PROD=nucleoporin 98kD /DB_XREF=gi:11120677 /UG=Hs.112255 nucleoporin 98kD /FL=gb:NM_005387.2 gb:AF071076.1	NM_005387		NP_624358
203196_at	0.00884	ATP-binding cassette, sub-family C (CFTR/MRP), member 4	AI948503	Hs.387834	NP_005836

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203197_s_at	0.038535	carnitine palmitoyltransferase II	AW157077	Hs.274336	NP_060357
203202_at	0.001116	glioma pathogenesis-related protein	AI950314	Hs.154762	NP_008974
203203_s_at	2.17E-04	gb:NM_007043.1 /DEF=Homo sapiens HIV-1 rev binding protein 2 (HRB2), mRNA. /FEA=mRNA /GEN=HRB2 /PROD=HIV-1 rev binding protein 2 /DB_XREF=gi:5902047 /UG=Hs.154762 HIV-1 rev binding protein 2 /FL=gb:U55766.1 gb:NM_007043.1	NM_007043		NP_008974
203204_s_at	0.007581	gb:BC002558.1 /DEF=Homo sapiens, KIAA0677 gene product, clone MGC:1972, mRNA, complete cds. /FEA=mRNA /PROD=KIAA0677 gene product /DB_XREF=gi:12803466 /UG=Hs.155983 KIAA0677 gene product /FL=gb:BC002558.1 gb:AB014577.1 gb:NM_014663.1	BC002558		NP_055478
203206_at	5.03E-04	gb:NM_014661.1 /DEF=Homo sapiens KIAA0140 gene product (KIAA0140), mRNA. /FEA=mRNA /GEN=KIAA0140 /PROD=KIAA0140 gene product /DB_XREF=gi:7661937 /UG=Hs.156016 KIAA0140 gene product /FL=gb:D50930.1 gb:NM_014661.1	NM_014661		NP_055476
203211_s_at	0.038535	Consensus includes gb:AK027038.1 /DEF=Homo sapiens cDNA: FLJ23385 fis, clone HEP16802. /FEA=mRNA /DB_XREF=gi:10440053 /UG=Hs.181326 KIAA1073 protein /FL=gb:AB028996.1 gb:NM_016156.1	AK027038		NP_057240
203213_at	0.034317	cell division cycle 2, G1 to S and G2 to M	AL524035	Hs.334562	NP_203698
203218_at	0.026891	mitogen-activated protein kinase 9	W37431	Hs.246857	NP_620709

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203219_s_at	0.035763	gb:NM_000485.1 /DEF=Homo sapiens adenine phosphoribosyltransferase (APRT), mRNA. /FEA=mRNA /GEN=APRT /PROD=adenine phosphoribosyltransferase /DB_XREF=gi:4502170 /UG=Hs.28914 adenine phosphoribosyltransferase /FL=gb:NM_000485.1	NM_000485		NP_000476
203223_at	0.034317	gb:NM_004703.1 /DEF=Homo sapiens rabaptin-5 (RAB5EP), mRNA. /FEA=mRNA /GEN=RAB5EP /PROD=rabaptin-5 /DB_XREF=gi:4759005 /UG=Hs.326056 rabaptin-5 /FL=gb:NM_004703.1	NM_004703		NP_004694
203232_s_at	2.05E-04	gb:NM_000332.1 /DEF=Homo sapiens spinocerebellar ataxia 1 (olivopontocerebellar ataxia 1, autosomal dominant, ataxin 1) (SCA1), mRNA. /FEA=mRNA /GEN=SCA1 /PROD=ataxin 1 /DB_XREF=gi:4506792 /UG=Hs.74520 spinocerebellar ataxia 1 (olivopontocerebellar ataxia 1, autosomal dominant, ataxin 1) /FL=gb:NM_000332.1	NM_000332		
203234_at	7.31E-04	gb:NM_003364.1 /DEF=Homo sapiens uridine phosphorylase (UP), mRNA. /FEA=mRNA /GEN=UP /PROD=uridine phosphorylase /DB_XREF=gi:4507838 /UG=Hs.77573 uridine phosphorylase /FL=gb:BC001405.1 gb:NM_003364.1	NM_003364		NP_003355

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203243_s_at	3.26E-04	gb:NM_006457.1 /DEF=Homo sapiens LIM protein (similar to rat protein kinase C-binding enigma) (LIM), mRNA. /FEA=mRNA /GEN=LIM /PROD=LIM protein (similar to rat protein kinaseC-binding enigma) /DB_XREF=gi:5453713 /UG=Hs.154103 LIM protein (similar to rat protein kinase C-binding enigma) /FL=gb:AF061258.1 gb:NM_006457.1	NM_006457		NP_006448
203244_at	0.021448	gb:NM_000319.1 /DEF=Homo sapiens peroxisome receptor 1 (PXR1), mRNA. /FEA=mRNA /GEN=PXR1 /PROD=peroxisome receptor 1 /DB_XREF=gi:4506346 /UG=Hs.158084 peroxisome receptor 1 /FL=gb:NM_000319.1 gb:U19721.1	NM_000319		NP_000310
203247_s_at	0.001618	gb:BC003566.1 /DEF=Homo sapiens, zinc finger protein 24 (KOX 17), clone MGC:2057, mRNA, complete cds. /FEA=mRNA /PROD=zinc finger protein 24 (KOX 17) /DB_XREF=gi:13097725 /UG=Hs.183593 zinc finger protein 24 (KOX 17) /FL=gb:BC003566.1 gb:U68536.1 gb:AF038964.1 gb:NM_006965.1	BC003566		NP_008896
203249_at	0.041055	gb:AB002386.1 /DEF=Human mRNA for KIAA0388 gene, complete cds. /FEA=mRNA /GEN=KIAA0388 /DB_XREF=gi:2224716 /UG=Hs.194669 enhancer of zeste (Drosophila) homolog 1 /FL=gb:U50315.1 gb:AB002386.1 gb:NM_001991.1	AB002386		NP_001982
203253_s_at	0.001116	gb:NM_015216.1 /DEF=Homo sapiens KIAA0433 protein (KIAA0433), mRNA. /FEA=mRNA /GEN=KIAA0433 /PROD=KIAA0433 protein /DB_XREF=gi:7662117 /UG=Hs.26179 KIAA0433 protein /FL=gb:NM_015216.1	NM_015216		NP_056031

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203258_at	0.033381	gb:NM_006442.1 /DEF=Homo sapiens DR1-associated protein 1 (negative cofactor 2 alpha) (DRAP1), mRNA. /FEA=mRNA /GEN=DRAP1 /PROD=DR1-associated protein 1 (negative cofactor 2alpha) /DB_XREF=gi:5453635 /UG=Hs.295362 DR1-associated protein 1 (negative cofactor 2 alpha) /FL=gb:U41843.1 gb:NM_006442.1	NM_006442		NP_006433
203261_at	0.02145	gb:NM_006571.1 /DEF=Homo sapiens novel RGD-containing protein (WS-3), mRNA. /FEA=mRNA /GEN=WS-3 /PROD=novel RGD-containing protein /DB_XREF=gi:5730115 /UG=Hs.39913 novel RGD-containing protein /FL=gb:D84145.1 gb:NM_006571.1	NM_006571		NP_006562
203265_s_at	1.26E-04	mitogen-activated protein kinase kinase 4	AA810268	Hs.75217	
203266_s_at	2.05E-04	gb:NM_003010.1 /DEF=Homo sapiens mitogen-activated protein kinase kinase 4 (MAP2K4), mRNA. /FEA=mRNA /GEN=MAP2K4 /PROD=mitogen-activated protein kinase kinase 4 /DB_XREF=gi:4506888 /UG=Hs.75217 mitogen-activated protein kinase kinase 4 /FL=gb:NM_003010.1 gb:L36870.1 gb:U17743.1	NM_003010		NP_003001
203272_s_at	0.004733	lung cancer candidate	BF308548	Hs.8186	NP_009206
203273_s_at	0.003368	gb:NM_007275.1 /DEF=Homo sapiens lung cancer candidate (FUS1), mRNA. /FEA=mRNA /GEN=FUS1 /PROD=lung cancer candidate /DB_XREF=gi:6005759 /UG=Hs.8186 lung cancer candidate /FL=gb:AF055479.1 gb:NM_007275.1	NM_007275		NP_009206

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203274_at	1.26E-04	gb:NM_012151.2 /DEF=Homo sapiens coagulation factor VIII-associated (intronic transcript) (F8A), mRNA. /FEA=mRNA /GEN=F8A /PROD=coagulation factor VIII-associated protein /DB_XREF=gi:12056462 /UG=Hs.83363 coagulation factor VIII-associated (intronic transcript) /FL=gb:NM_012151.2	NM_012151		NP_036283
203275_at	0.00228	gb:NM_002199.2 /DEF=Homo sapiens interferon regulatory factor 2 (IRF2), mRNA. /FEA=mRNA /GEN=IRF2 /PROD=interferon regulatory factor 2 /DB_XREF=gi:4755144 /UG=Hs.83795 interferon regulatory factor 2 /FL=gb:NM_002199.2	NM_002199		NP_002190
203276_at	0.026891	gb:NM_005573.1 /DEF=Homo sapiens lamin B1 (LMNB1), mRNA. /FEA=mRNA /GEN=LMNB1 /PROD=lamin B1 /DB_XREF=gi:5031876 /UG=Hs.89497 lamin B1 /FL=gb:M34458.1 gb:NM_005573.1	NM_005573		NP_005564
203279_at	3.26E-04	gb:NM_014674.1 /DEF=Homo sapiens KIAA0212 gene product (KIAA0212), mRNA. /FEA=mRNA /GEN=KIAA0212 /PROD=KIAA0212 gene product /DB_XREF=gi:7662001 /UG=Hs.154332 KIAA0212 gene product /FL=gb:D86967.1 gb:NM_014674.1	NM_014674		NP_055489

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203282_at	1.26E-04	gb:NM_000158.1 /DEF=Homo sapiens glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA. /FEA=mRNA /GEN=GBE1 /PROD=glucan (1,4-alpha-), branching enzyme 1(glycogen branching enzyme) /DB_XREF=gi:4557618 /UG=Hs.1691 glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) /FL=gb:L07956.1 gb:NM_000158.1	NM_000158		NP_000149
203283_s_at	0.009682	Consensus includes gb:AK023260.1 /DEF=Homo sapiens cDNA FLJ13198 fis, clone NT2RP3004454, highly similar to Homo sapiens mRNA for KIAA0448 protein. /FEA=mRNA /DB_XREF=gi:10435114 /UG=Hs.169939 heparan sulfate 2-O-sulfotransferase /FL=gb:NM_012262.2 gb:AB007917.1 gb:AB024568.1	NM_012262		NP_036394
203284_s_at	3.26E-04	Consensus includes gb:AW151887 /FEA=EST /DB_XREF=gi:6199872 /DB_XREF=est:xf72c12.x1 /CLONE=IMAGE:2623606 /UG=Hs.169939 heparan sulfate 2-O-sulfotransferase /FL=gb:NM_012262.2 gb:AB007917.1 gb:AB024568.1	NM_012262		NP_036394
203285_s_at	1.26E-04	gb:NM_012262.2 /DEF=Homo sapiens heparan sulfate 2-O-sulfotransferase (HS2ST1), mRNA. /FEA=mRNA /GEN=HS2ST1 /PROD=heparan sulfate 2-O-sulfotransferase 1 /DB_XREF=gi:12545388 /UG=Hs.169939 heparan sulfate 2-O-sulfotransferase /FL=gb:NM_012262.2 gb:AB007917.1 gb:AB024568.1	NM_012262		NP_036394

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Prot in Accession No.
203286_at	2.05E-04	gb:NM_014901.1 /DEF=Homo sapiens KIAA1100 protein (KIAA1100), mRNA. /FEA=mRNA /GEN=KIAA1100 /PROD=KIAA1100 protein /DB_XREF=gi:7662485 /UG=Hs.179946 KIAA1100 protein /FL=gb:AB029023.1 gb:NM_014901.1	NM_014901		NP_055716
203288_at	0.003529	gb:NM_014686.1 /DEF=Homo sapiens KIAA0355 gene product (KIAA0355), mRNA. /FEA=mRNA /GEN=KIAA0355 /PROD=KIAA0355 gene product /DB_XREF=gi:7662075 /UG=Hs.186840 KIAA0355 gene product /FL=gb:AB002353.1 gb:NM_014686.1	NM_014686		NP_055501
203291_at	5.03E-04	gb:NM_013316.1 /DEF=Homo sapiens CCR4-NOT transcription complex, subunit 4 (CNOT4), mRNA. /FEA=mRNA /GEN=CNOT4 /PROD=CCR4-NOT transcription complex, subunit 4 /DB_XREF=gi:7019466 /UG=Hs.20423 CCR4-NOT transcription complex, subunit 4 /FL=gb:U71267.1 gb:NM_013316.1	NM_013316		NP_037448
203292_s_at	0.001116	gb:NM_021729.2 /DEF=Homo sapiens vacuolar protein sorting 11 (yeast homolog) (VPS11), mRNA. /FEA=mRNA /GEN=VPS11 /PROD=vacuolar protein sorting 11 (yeast homolog) /DB_XREF=gi:11141864 /UG=Hs.234282 vacuolar protein sorting 11 (yeast homolog) /FL=gb:AB027508.2 gb:NM_021729.2 gb:AF308800.1	NM_021729		NP_068375
203298_s_at	0.019761	gb:NM_004973.2 /DEF=Homo sapiens jumonji (mouse) homolog (JMJ), mRNA. /FEA=mRNA /GEN=JMJ /PROD=jumonji /DB_XREF=gi:11863151 /UG=Hs.40154 jumonji (mouse) homolog /FL=gb:NM_004973.2 gb:U57592.1	NM_004973		NP_004964

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203300_x_at	0.041055	gb:NM_003916.1 /DEF=Homo sapiens adaptor-related protein complex 1, sigma 2 subunit (AP1S2), mRNA. /FEA=mRNA /GEN=AP1S2 /PROD=adaptor-related protein complex 1, sigma 2 subunit /DB_XREF=gi:4506956 /UG=Hs.40368 adaptor-related protein complex 1, sigma 2 subunit /FL=gb:AF251295.1 gb:BC001117.1 gb:AB015320.1 gb:NM_003916.1	NM_003916		NP_003907
203304_at	0.007942	gb:NM_012342.1 /DEF=Homo sapiens putative transmembrane protein (NMA), mRNA. /FEA=mRNA /GEN=NMA /PROD=putative transmembrane protein /DB_XREF=gi:6912533 /UG=Hs.78776 putative transmembrane protein /FL=gb:U23070.1 gb:NM_012342.1	NM_012342		NP_036474
203305_at	0.013223	gb:NM_000129.2 /DEF=Homo sapiens coagulation factor XIII, A1 polypeptide (F13A1), mRNA. /FEA=mRNA /GEN=F13A1 /PROD=coagulation factor XIII A1 subunit precursor /DB_XREF=gi:9961355 /UG=Hs.80424 coagulation factor XIII, A1 polypeptide /FL=gb:M14354.1 gb:NM_000129.2	NM_000129		NP_000120
203306_s_at	0.013223	gb:NM_006416.1 /DEF=Homo sapiens solute carrier family 35 (CMP-sialic acid transporter), member 1 (SLC35A1), mRNA. /FEA=mRNA /GEN=SLC35A1 /PROD=solute carrier family 35 (CMP-sialic acid transporter), member 1 /DB_XREF=gi:5453620 /UG=Hs.82921 solute carrier family 35 (CMP-sialic acid transporter), member 1 /FL=gb:D87969.1 gb:NM_006416.1	NM_006416		NP_006407

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203309_s_at	0.003171	gb:NM_000195.1 /DEF=Homo sapiens Hermansky-Pudlak syndrome (HPS), mRNA. /FEA=mRNA /GEN=HPS /PROD=Hermansky-Pudlak syndrome protein /DB_XREF=gi:4504484 /UG=Hs.83951 Hermansky-Pudlak syndrome /FL=gb:U65676.1 gb:NM_000195.1	NM_000195		NP_000186
203310_at	0.004341	gb:NM_007269.1 /DEF=Homo sapiens syntaxin binding protein 3 (STXBP3), mRNA. /FEA=mRNA /GEN=STXBP3 /PROD=syntaxin 4 binding protein /DB_XREF=gi:6005885 /UG=Hs.8813 syntaxin binding protein 3 /FL=gb:D63506.1 gb:AF032922.1 gb:NM_007269.1	NM_007269		NP_009200
203318_s_at	1.26E-04	gb:NM_021964.1 /DEF=Homo sapiens zinc finger protein 148 (pHZ-52) (ZNF148), mRNA. /FEA=mRNA /GEN=ZNF148 /PROD=zinc finger protein 148 (pHZ-52) /DB_XREF=gi:11415035 /UG=Hs.112180 zinc finger protein 148 (pHZ-52) /FL=gb:NM_021964.1 gb:L04282.1	NM_021964		NP_068799
203319_s_at	7.58E-04	gb:L04282.1 /DEF=Human CACCC box-binding protein mRNA, complete cds. /FEA=mRNA /PROD=CACCC box-binding protein /DB_XREF=gi:388318 /UG=Hs.112180 zinc finger protein 148 (pHZ-52) /FL=gb:NM_021964.1 gb:L04282.1	L04282		NP_068799
203320_at	3.26E-04	gb:NM_005475.1 /DEF=Homo sapiens lymphocyte adaptor protein (LNK), mRNA. /FEA=mRNA /GEN=LNK /PROD=lymphocyte adaptor protein /DB_XREF=gi:4885454 /UG=Hs.13131 lymphocyte adaptor protein /FL=gb:AF055581.1 gb:NM_005475.1	NM_005475		NP_005466

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203321_s_at	3.26E-04	Consensus includes gb:AK022688.1 /DEF=Homo sapiens cDNA FLJ12626 fis, clone NT2RM4001810, weakly similar to AGGRECAN CORE PROTEIN PRECURSOR. /FEA=CDS /DB_XREF=gi:10434228 /UG=Hs.131915 KIAA0863 protein /FL=gb:AB020670.1 gb:NM_014913.1	NM_014913		NP_055728
203328_x_at	0.012168	gb:NM_004969.1 /DEF=Homo sapiens insulin-degrading enzyme (IDE), mRNA. /FEA=mRNA /GEN=IDE /PROD=insulysin /DB_XREF=gi:4826769 /UG=Hs.1508 insulin-degrading enzyme /FL=gb:M21188.1 gb:NM_004969.1	NM_004969		NP_004960
203331_s_at	0.020576	gb:U53470.1 /DEF=Human signaling inositol polyphosphate phosphatase SHIP II mRNA, complete cds. /FEA=mRNA /PROD=signaling inositol polyphosphate phosphataseSHIP II /DB_XREF=gi:4097284 /UG=Hs.155939 inositol polyphosphate-5-phosphatase, 145kD /FL=gb:U50040.1 gb:U57650.1 gb:U84400.1 gb:U53470.1 gb:NM_005541.1	U53470		NP_005532
203334_at	0.025245	gb:NM_004941.1 /DEF=Homo sapiens DEADH (Asp-Glu-Ala-AspHis) box polypeptide 8 (RNA helicase) (DDX8), mRNA. /FEA=mRNA /GEN=DDX8 /PROD=DEADH (Asp-Glu-Ala-AspHis) box polypeptide 8 /DB_XREF=gi:4826689 /UG=Hs.171872 DEADH (Asp-Glu-Ala-AspHis) box polypeptide 8 (RNA helicase) /FL=gb:D50487.1 gb:NM_004941.1	NM_004941		NP_004932

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203340_s_at	0.007943	gb:NM_003705.1 /DEF=Homo sapiens solute carrier family 25 (mitochondrial carrier, Aralar), member 12 (SLC25A12), mRNA. /FEA=mRNA /GEN=SLC25A12 /PROD=solute carrier family 25 (mitochondrial carrier, Aralar), member 12 /DB_XREF=gi:4507006 /UG=Hs.179866 solute carrier family 25 (mitochondrial carrier, Aralar), member 12 /FL=gb:NM_003705.1	NM_003705		NP_003696
203342_at	0.003741	gb:NM_005834.1 /DEF=Homo sapiens translocase of inner mitochondrial membrane 17 (yeast) homolog B (TIM17B), mRNA. /FEA=mRNA /GEN=TIM17B /PROD=translocase of inner mitochondrial membrane 17(yeast) homolog B /DB_XREF=gi:5032180 /UG=Hs.19105 translocase of inner mitochondrial membrane 17 (yeast) homolog B /FL=gb:AJ005895.1 gb:AF034790.1 gb:AF077039.1 gb:NM_005834.1	NM_005834		NP_005825
203346_s_at	0.030771	gb:AF072814.1 /DEF=Homo sapiens PHD finger DNA binding protein isoform 1 (M96) mRNA, alternatively spliced, complete cds. /FEA=mRNA /GEN=M96 /PROD=PHD finger DNA binding protein isoform 1 /DB_XREF=gi:3342451 /UG=Hs.31016 putative DNA binding protein /FL=gb:AF072814.1 gb:NM_007358.1	AF072814		NP_031384

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203347_s_at	0.010205	gb:NM_007358.1 /DEF=Homo sapiens putative DNA binding protein (M96), mRNA. /FEA=mRNA /GEN=M96 /PROD=putative DNA binding protein /DB_XREF=gi:6678763 /UG=Hs.31016 putative DNA binding protein /FL=gb:AF072814.1 gb:NM_007358.1	NM_007358		NP_031384
203350_at	0.025464	gb:NM_001128.1 /DEF=Homo sapiens adaptor-related protein complex 1, gamma 1 subunit (AP1G1), mRNA. /FEA=mRNA /GEN=AP1G1 /PROD=adaptor-related protein complex 1, gamma 1 subunit /DB_XREF=gi:4501978 /UG=Hs.5344 adaptor-related protein complex 1, gamma 1 subunit /FL=gb:AB015317.1 gb:NM_001128.1	NM_001128		NP_001119
203353_s_at	0.005039	gb:NM_015846.1 /DEF=Homo sapiens methyl-CpG binding domain protein 1 (MBD1), transcript variant 1, mRNA. /FEA=mRNA /GEN=MBD1 /PROD=methyl-CpG binding domain protein 1, isoform 1 /DB_XREF=gi:7710138 /UG=Hs.6211 methyl-CpG binding domain protein 1 /FL=gb:AF078830.1 gb:NM_015846.1	NM_015846		NP_056723
203359_s_at	1.26E-04	c-myc binding protein	AL525412	Hs.78221	NP_036465
203360_s_at	1.26E-04	gb:D50692.1 /DEF=Homo sapiens mRNA for c-myc binding protein, complete cds. /FEA=mRNA /PROD=c-myc binding protein /DB_XREF=gi:1785850 /UG=Hs.78221 c-myc binding protein /FL=gb:D50692.1 gb:AB007191.2 gb:NM_012333.2	D50692		NP_036465

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203361_s_at	0.011725	gb:NM_012333.2 /DEF=Homo sapiens c-myc binding protein (MYCBP), mRNA. /FEA=mRNA /GEN=MYCBP /PROD=c-myc binding protein /DB_XREF=gi:8850230 /UG=Hs.78221 c-myc binding protein /FL=gb:D50692.1 gb:AB007191.2 gb:NM_012333.2	NM_012333		NP_036465
203362_s_at	7.58E-04	gb:NM_002358.2 /DEF=Homo sapiens MAD2 (mitotic arrest deficient, yeast, homolog)-like 1 (MAD2L1), mRNA. /FEA=mRNA /GEN=MAD2L1 /PROD=MAD2-like 1 /DB_XREF=gi:6466452 /UG=Hs.79078 MAD2 (mitotic arrest deficient, yeast, homolog)-like 1 /FL=gb:BC000356.1 gb:U65410.1 gb:NM_002358.2 gb:U31278.1	NM_002358		NP_002349
203370_s_at	0.033381	gb:NM_005451.2 /DEF=Homo sapiens enigma (LIM domain protein) (ENIGMA), mRNA. /FEA=mRNA /GEN=ENIGMA /PROD=enigma protein /DB_XREF=gi:11496884 /UG=Hs.102948 enigma (LIM domain protein) /FL=gb:NM_005451.2 gb:BC001093.1 gb:AF265209.1	NM_005451		NP_005442
203371_s_at	7.58E-04	gb:NM_002491.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3 (12kD, B12) (NDUFB3), mRNA. /FEA=mRNA /GEN=NDUFB3 /PROD=NADH dehydrogenase (ubiquinone) 1 betasubcomplex, 3 (12kD, B12) /DB_XREF=gi:4505360 /UG=Hs.109760 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3 (12kD, B12) /FL=gb:AF047183.1 gb:NM_002491.1	NM_002491		NP_002482

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203372_s_at	0.048741	gb:AB004903.1 /DEF=Homo sapiens mRNA for STAT induced STAT inhibitor-2, complete cds. /FEA=mRNA /PROD=STAT induced STAT inhibitor-2 /DB_XREF=gi:2443360 /UG=Hs.110776 STAT induced STAT inhibitor-2 /FL=gb:AB004903.1 gb:AB006966.1 gb:AF037989.1 gb:AF020590.1 gb:NM_003877.1	AB004903		NP_003868
203376_at	0.007779	pre-mRNA splicing factor 17	BG528818	Hs.116674	NP_056975
203379_at	5.03E-04	gb:NM_002953.1 /DEF=Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 1 (RPS6KA1), mRNA. /FEA=mRNA /GEN=RPS6KA1 /PROD=ribosomal protein S6 kinase, 90kD, polypeptide1 /DB_XREF=gi:4506732 /UG=Hs.149957 ribosomal protein S6 kinase, 90kD, polypeptide 1 /FL=gb:L07597.1 gb:NM_002953.1	NM_002953		NP_002944
203384_s_at	0.005852	gb:NM_002077.1 /DEF=Homo sapiens golgi autoantigen, golgin subfamily a, 1 (GOLGA1), mRNA. /FEA=mRNA /GEN=GOLGA1 /PROD=golgin 97 /DB_XREF=gi:4504062 /UG=Hs.172647 golgi autoantigen, golgin subfamily a, 1 /FL=gb:U51587.1 gb:NM_002077.1	NM_002077		NP_002068
203385_at	0.007779	gb:NM_001345.1 /DEF=Homo sapiens diacylglycerol kinase, alpha (80kD) (DGKA), mRNA. /FEA=mRNA /GEN=DGKA /PROD=diacylglycerol kinase, alpha (80kD) /DB_XREF=gi:11415023 /UG=Hs.172690 diacylglycerol kinase, alpha (80kD) /FL=gb:NM_001345.1 gb:AF064770.1	NM_001345		NP_001336

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203387_s_at	0.033381	gb:NM_014832.1 /DEF=Homo sapiens KIAA0603 gene product (KIAA0603), mRNA. /FEA=mRNA /GEN=KIAA0603 /PROD=KIAA0603 gene product /DB_XREF=gi:7662197 /UG=Hs.173802 KIAA0603 gene product /FL=gb:AB011175.1 gb:NM_014832.1	NM_014832		NP_055647
203394_s_at	0.00423	hairy homolog (Drosophila)	BE973687	Hs.250666	NP_005515
203397_s_at	0.003171	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3)	BF063271	Hs.278611	NP_004473
203406_at	0.004341	gb:NM_005926.1 /DEF=Homo sapiens microfibrillar-associated protein 1 (MFAP1), mRNA. /FEA=mRNA /GEN=MFAP1 /PROD=microfibrillar-associated protein 1 /DB_XREF=gi:5174552 /UG=Hs.61418 microfibrillar-associated protein 1 /FL=gb:U04209.1 gb:NM_005926.1	NM_005926		NP_005917
203408_s_at	3.26E-04	gb:NM_002971.1 /DEF=Homo sapiens special AT-rich sequence binding protein 1 (binds to nuclear matrixscaffold-associating DNAs) (SATB1), mRNA. /FEA=mRNA /GEN=SATB1 /PROD=special AT-rich sequence binding protein 1(binds to nuclear matrixscaffold-associating DNAs) /DB_XREF=gi:4506790 /UG=Hs.74592 special AT-rich sequence binding protein 1 (binds to nuclear matrixscaffold-associating DNAs) /FL=gb:M97287.1 gb:NM_002971.1	NM_002971		NP_002962
203415_at	1.26E-04	gb:NM_013232.1 /DEF=Homo sapiens programmed cell death 6 (PDCD6), mRNA. /FEA=mRNA /GEN=PDCD6 /PROD=programmed cell death 6 /DB_XREF=gi:7019484 /UG=Hs.80019 programmed cell death 6 /FL=gb:AF035606.1 gb:U58773.1 gb:NM_013232.1	NM_013232		NP_037364

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203416_at	0.00228	gb:NM_000560.1 /DEF=Homo sapiens CD53 antigen (CD53), mRNA. /FEA=mRNA /GEN=CD53 /PROD=CD53 antigen /DB_XREF=gi:10834971 /UG=Hs.82212 CD53 antigen /FL=gb:NM_000560.1 gb:M60871.1 gb:M37033.1	NM_000560		NP_000551
203420_at	0.033381	gb:NM_016255.1 /DEF=Homo sapiens Autosomal Highly Conserved Protein (AHCP), mRNA. /FEA=mRNA /GEN=AHCP /PROD=Autosomal Highly Conserved Protein /DB_XREF=gi:7705267 /UG=Hs.95260 Autosomal Highly Conserved Protein /FL=gb:AF097027.1 gb:NM_016255.1	NM_016255		NP_057339
203426_s_at	0.01659	gb:M65062.1 /DEF=Human insulin-like growth factor binding protein 5 (IGFBP-5) mRNA, complete cds. /FEA=mRNA /GEN=IGFBP5 /PROD=insulin-like growth factor binding protein 5 /DB_XREF=gi:184819 /UG=Hs.107169 insulin-like growth factor binding protein 5 /FL=gb:NM_000599.1 gb:M62782.1 gb:M65062.1 gb:AF055033.1	M65062		NP_000590
203430_at	1.26E-04	gb:NM_014320.1 /DEF=Homo sapiens putative heme-binding protein (SOUL), mRNA. /FEA=mRNA /GEN=SOUL /PROD=putative heme-binding protein /DB_XREF=gi:7657602 /UG=Hs.111029 putative heme-binding protein /FL=gb:AF117616.1 gb:NM_014320.1	NM_014320		NP_055135
203432_at	0.02145	thymopoietin	AW272611	Hs.11355	NP_003267

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203433_at	0.00161	gb:NM_006441.1 /DEF=Homo sapiens 5,10-methenyltetrahydrofolate synthetase (5-formyltetrahydrofolate cyclo-ligase) (MTHFS), mRNA. /FEA=mRNA /GEN=MTHFS /PROD=5,10-methenyltetrahydrofolate synthetase(5-formyltetrahydrofolate cyclo-ligase) /DB_XREF=gi:5453745 /UG=Hs.118131 5,10-methenyltetrahydrofolate synthetase (5-formyltetrahydrofolate cyclo-ligase) /FL=gb:NM_006441.1 gb:L38928.1	NM_006441		NP_006432
203435_s_at	0.027243	gb:NM_007287.1 /DEF=Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10) (MME), transcript variant 1bis, mRNA. /FEA=mRNA /GEN=MME /PROD=membrane metallo-endopeptidase /DB_XREF=gi:6042199 /UG=Hs.1298 membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10) /FL=gb:J03779.1 gb:NM_007287.1 gb:NM_007288.1	NM_007287		NP_009220
203437_at	1.26E-04	gb:NM_003876.1 /DEF=Homo sapiens putative receptor protein (PMI), mRNA. /FEA=mRNA /GEN=PMI /PROD=putative receptor protein /DB_XREF=gi:4505900 /UG=Hs.15196 putative receptor protein /FL=gb:BC002819.1 gb:BC005268.1 gb:NM_003876.1	NM_003876		NP_003867

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203445_s_at	0.00161	gb:NM_005730.1 /DEF=Homo sapiens conserved gene amplified in osteosarcoma (OS4), mRNA. /FEA=mRNA /GEN=OS4 /PROD=conserved gene amplified in osteosarcoma /DB_XREF=gi:5031964 /UG=Hs.180669 conserved gene amplified in osteosarcoma /FL=gb:U81556.1 gb:NM_005730.1	NM_005730		NP_005721
203447_at	0.013223	proteasome (prosome, macropain) 26S subunit, non-ATPase, 5	AU157008	Hs.273721	NP_005038
203457_at	2.05E-04	gb:NM_003569.1 /DEF=Homo sapiens syntaxin 7 (STX7), mRNA. /FEA=mRNA /GEN=STX7 • /PROD=syntaxin 7 /DB_XREF=gi:4507294 /UG=Hs.299166 syntaxin 7 /FL=gb:U77942.1 gb:NM_003569.1	NM_003569		NP_003560
203459_s_at	0.026891	gb:NM_022575.1 /DEF=Homo sapiens vacuolar protein sorting 16 (yeast homolog) (VPS16), mRNA. /FEA=mRNA /GEN=VPS16 /PROD=vacuolar protein sorting 16 (yeast homolog) /DB_XREF=gi:12007657 /UG=Hs.302441 vacuolar protein sorting 16 (yeast homolog) /FL=gb:AF308801.1 gb:NM_022575.1	NM_022575		NP_536339
203460_s_at	1.26E-04	gb:NM_007318.1 /DEF=Homo sapiens presenilin 1 (Alzheimer disease 3) (PSEN1), transcript variant I-463, mRNA. /FEA=mRNA /GEN=PSEN1 /PROD=presenilin 1 isoform I-463 /DB_XREF=gi:7549812 /UG=Hs.3260 presenilin 1 (Alzheimer disease 3) /FL=gb:U40379.1 gb:L76517.1 gb:NM_007318.1	NM_007318		NP_015558

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203471_s_at	7.58E-04	gb:NM_002664.1 /DEF=Homo sapiens pleckstrin (PLEK), mRNA. /FEA=mRNA /GEN=PLEK /PROD=pleckstrin /DB_XREF=gi:4505878 /UG=Hs.77436 pleckstrin /FL=gb:NM_002664.1	NM_002664		NP_002655
203478_at	0.004341	gb:NM_002494.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (6kD, KFYI) (NDUFC1), mRNA. /FEA=mRNA /GEN=NDUFC1 /PROD=NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (6kD, KFYI) /DB_XREF=gi:4505366 /UG=Hs.84549 NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (6kD, KFYI) /FL=gb:AF047184.1 gb:AF047435.1 gb:NM_002494.1	NM_002494		NP_002485
203484_at	0.004341	gb:NM_014302.1 /DEF=Homo sapiens Sec61 gamma (SEC61G), mRNA. /FEA=mRNA /GEN=SEC61G /PROD=Sec61 gamma /DB_XREF=gi:7657545 /UG=Hs.9950 Sec61 gamma /FL=gb:AF054184.1 gb:NM_014302.1	NM_014302		NP_055117
203486_s_at	2.05E-04	DKFZP434A043 protein	BF195973	Hs.102708	NP_056211
203487_s_at	3.26E-04	gb:NM_015396.1 /DEF=Homo sapiens DKFZP434A043 protein (DKFZP434A043), mRNA. /FEA=mRNA /GEN=DKFZP434A043 /PROD=DKFZP434A043 protein /DB_XREF=gi:7661561 /UG=Hs.102708 DKFZP434A043 protein /FL=gb:NM_015396.1	NM_015396		NP_056211
203497_at	5.03E-04	gb:NM_004774.1 /DEF=Homo sapiens PPAR binding protein (PPARBP), mRNA. /FEA=mRNA /GEN=PPARBP /PROD=thyroid hormone receptor interactor 2 /DB_XREF=gi:4759265 /UG=Hs.15589 PPAR binding protein /FL=gb:AF055994.1 gb:NM_004774.1 gb:AF283812.1	NM_004774		NP_004765

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203501_at	3.26E-04	gb:NM_006102.1 /DEF=Homo sapiens plasma glutamate carboxypeptidase (PGCP), mRNA. /FEA=mRNA /GEN=PGCP /PROD=plasma glutamate carboxypeptidase /DB_XREF=gi:5174626 /UG=Hs.197335 plasma glutamate carboxypeptidase /FL=gb:AF119386.1 gb:NM_006102.1	NM_006102		
203502_at	1.26E-04	gb:NM_001724.1 /DEF=Homo sapiens 2,3-bisphosphoglycerate mutase (BPGM), mRNA. /FEA=mRNA /GEN=BPGM /PROD=2,3-bisphosphoglycerate mutase /DB_XREF=gi:4502444 /UG=Hs.198365 2,3-bisphosphoglycerate mutase /FL=gb:NM_001724.1	NM_001724		NP_001715
203506_s_at	6.68E-04	gb:NM_005120.1 /DEF=Homo sapiens trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit) (TNRC11), mRNA. /FEA=mRNA /GEN=TNRC11 /PROD=trinucleotide repeat containing 11(THR-associated protein, 230 kDa subunit) /DB_XREF=gi:4827041 /UG=Hs.211607 trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit) /FL=gb:U80742.1 gb:AF071309.1 gb:AF117755.1 gb:NM_005120.1	NM_005120		NP_005111
203507_at	5.96E-04	gb:NM_001251.1 /DEF=Homo sapiens CD68 antigen (CD68), mRNA. /FEA=mRNA /GEN=CD68 /PROD=CD68 antigen /DB_XREF=gi:4557434 /UG=Hs.246381 CD68 antigen /FL=gb:NM_001251.1	NM_001251		NP_001242

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203508_at	2.05E-04	gb:NM_001066.1 /DEF=Homo sapiens tumor necrosis factor receptor superfamily, member 1B (TNFRSF1B), mRNA. /FEA=mRNA /GEN=TNFRSF1B /PROD=tumor necrosis factor receptor 2 (75kD) /DB_XREF=gi:4507576 /UG=Hs.256278 tumor necrosis factor receptor superfamily, member 1B /FL=gb:M32315.1 gb:M55994.1 gb:NM_001066.1	NM_001066		NP_001057
203511_s_at	5.03E-04	gb:AF041432.1 /DEF=Homo sapiens bet3 (BET3) mRNA, complete cds. /FEA=mRNA /GEN=BET3 /PROD=bet3 /DB_XREF=gi:2791803 /UG=Hs.288013 similar to yeast BET3 (S. cerevisiae) /FL=gb:AF041432.1 gb:NM_014408.1	AF041432		NP_055223
203512_at	7.58E-04	gb:NM_014408.1 /DEF=Homo sapiens similar to yeast BET3 (S. cerevisiae) (BET3), mRNA. /FEA=mRNA /GEN=BET3 /PROD=similar to yeast BET3 (S. cerevisiae) /DB_XREF=gi:7656925 /UG=Hs.288013 similar to yeast BET3 (S. cerevisiae) /FL=gb:AF041432.1 gb:NM_014408.1	NM_014408		NP_055223
203513_at	0.007779	gb:NM_025137.1 /DEF=Homo sapiens hypothetical protein FLJ21439 (FLJ21439), mRNA. /FEA=mRNA /GEN=FLJ21439 /PROD=hypothetical protein FLJ21439 /DB_XREF=gi:13376718 /UG=Hs.288872 hypothetical protein FLJ21439 /FL=gb:NM_025137.1	NM_025137		NP_079413
203514_at	0.010205	mitogen-activated protein kinase kinase kinase 3	BF971923	Hs.29282	NP_002392

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203518_at	0.001116	gb:NM_000081.1 /DEF=Homo sapiens Chediak-Higashi syndrome 1 (CHS1), mRNA. /FEA=mRNA /GEN=CHS1 /PROD=beige protein homolog /DB_XREF=gi:4502838 /UG=Hs.36508 Chediak-Higashi syndrome 1 /FL=gb:U67615.1 gb:NM_000081.1	NM_000081		NP_000072
203519_s_at	0.026891	gb:NM_015542.1 /DEF=Homo sapiens regulator of nonsense transcripts 2; DKFZP434D222 protein (RENT2), mRNA. /FEA=mRNA /GEN=RENT2 /PROD=regulator of nonsense transcripts 2 /DB_XREF=gi:11693131 /UG=Hs.3862 regulator of nonsense transcripts 2; DKFZP434D222 protein /FL=gb:AF301013.1 gb:NM_015542.1 gb:AY013249.1 gb:AF318574.1	NM_015542		NP_542166
203525_s_at	0.005852	adenomatosis polyposis coli	AI375486	Hs.75081	NP_000029
203526_s_at	0.033381	gb:M74088.1 /DEF=Human APC gene mRNA, complete cds. /FEA=mRNA /GEN=APC /DB_XREF=gi:182396 /UG=Hs.75081 adenomatosis polyposis coli /FL=gb:M74088.1 gb:M73548.1 gb:NM_000038.1	M74088		NP_000029
203529_at	7.58E-04	gb:NM_016294.1 /DEF=Homo sapiens serinethreonine protein phosphatase catalytic subunit (LOC51723), mRNA. /FEA=mRNA /GEN=LOC51723 /PROD=serinethreonine protein phosphatase catalyticsubunit /DB_XREF=gi:7706488 /UG=Hs.80324 serinethreonine protein phosphatase catalytic subunit /FL=gb:AF035158.2 gb:NM_016294.1	NM_016294		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203534_at	0.003171	gb:NM_014462.1 /DEF=Homo sapiens Lsm1 protein (LSM1), mRNA. /FEA=mRNA /GEN=LSM1 /PROD=Lsm1 protein /DB_XREF=gi:7657312 /UG=Hs.111783 Lsm1 protein /FL=gb:BC001767.1 gb:AF000177.1 gb:NM_014462.1	NM_014462		NP_055277
203544_s_at	3.26E-04	gb:NM_003473.1 /DEF=Homo sapiens signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 (STAM), mRNA. /FEA=mRNA /GEN=STAM /PROD=signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 /DB_XREF=gi:4507248 /UG=Hs.153487 signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 /FL=gb:U43899.1 gb:NM_003473.1	NM_003473		NP_003464
203559_s_at	0.020821	gb:NM_001091.1 /DEF=Homo sapiens amiloride binding protein 1 (amine oxidase (copper-containing)) (ABP1), mRNA. /FEA=mRNA /GEN=ABP1 /PROD=amiloride binding protein 1 precursor /DB_XREF=gi:4501850 /UG=Hs.75741 amiloride binding protein 1 (amine oxidase (copper-containing)) /FL=gb:M55602.1 gb:NM_001091.1 gb:U11862.1	NM_001091		NP_001082

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203560_at	1.32E-04	gb:NM_003878.1 /DEF=Homo sapiens gamma-glutamyl hydrolase (conjugase, foyl/polygammaglutamyl hydrolase) (GGH), mRNA. /FEA=mRNA /GEN=GGH /PROD=gamma-glutamyl hydrolase (conjugase, foyl/polygammaglutamyl hydrolase) precursor /DB_XREF=gi:4503986 /UG=Hs.78619 gamma-glutamyl hydrolase (conjugase, foyl/polygammaglutamyl hydrolase) /FL=gb:U55206.1 gb:NM_003878.1	NM_003878		NP_003869
203561_at	0.00228	gb:NM_021642.1 /DEF=Homo sapiens Fc fragment of IgG, low affinity IIa, receptor for (CD32) (FCGR2A), mRNA. /FEA=mRNA /GEN=FCGR2A /PROD=Fc fragment of IgG, low affinity IIa, receptor for (CD32) /DB_XREF=gi:11056051 /UG=Hs.78864 Fc fragment of IgG, low affinity IIa, receptor for (CD32) /FL=gb:NM_021642.1 gb:M31932.1 gb:J03619.1 gb:M28697.1	NM_021642		NP_067674
203566_s_at	0.00161	gb:NM_000645.1 /DEF=Homo sapiens amylo-1,6-glucosidase, 4-alpha-glucanotransferase (glycogen debranching enzyme, glycogen storage disease type III) (AGL), transcript variant 5, mRNA. /FEA=mRNA /GEN=AGL /PROD=amylo-1,6-glucosidase, 4-alpha-glucanotransferase isoform 2 /DB_XREF=gi:4557282 /UG=Hs.904 amylo-1,6-glucosidase, 4-alpha-glucanotransferase (glycogen debranching enzyme, glycogen storage disease type III) /FL=gb:M85168.1 gb:NM_000645.1	NM_000645		NP_000637

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203573_s_at	0.014715	gb:NM_004581.1 /DEF=Homo sapiens Rab geranylgeranyltransferase, alpha subunit (RABGGTA), mRNA. /FEA=mRNA /GEN=RABGGTA /PROD=Rab geranylgeranyltransferase, alpha subunit /DB_XREF=gi:4759015 /UG=Hs.78920 Rab geranylgeranyltransferase, alpha subunit /FL=gb:NM_004581.1	NM_004581		NP_004572
203574_at	5.03E-04	gb:NM_005384.1 /DEF=Homo sapiens nuclear factor, interleukin 3 regulated (NFIL3), mRNA. /FEA=mRNA /GEN=NFIL3 /PROD=nuclear factor, interleukin 3 regulated /DB_XREF=gi:4885516 /UG=Hs.79334 nuclear factor, interleukin 3 regulated /FL=gb:U26173.1 gb:NM_005384.1	NM_005384		NP_005375
203579_s_at	0.02145	Consensus includes gb:AI660619 /FEA=EST /DB_XREF=gi:4764202 /DB_XREF=est:wf21h04.x1 /CLONE=IMAGE:2351287 /UG=Hs.10315 solute carrier family 7 (cationic amino acid transporter, y+ system), member 6 /FL=gb:D87432.1 gb:NM_003983.1	NM_003983		NP_003974
203583_at	3.26E-04	gb:NM_014044.1 /DEF=Homo sapiens DKFZP564G0222 protein (DKFZP564G0222), mRNA. /FEA=mRNA /GEN=DKFZP564G0222 /PROD=DKFZP564G0222 protein /DB_XREF=gi:7661609 /UG=Hs.13370 DKFZP564G0222 protein /FL=gb:AF077038.1 gb:AL080115.1 gb:NM_014044.1	NM_014044		NP_054763

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203584_at	0.010205	gb:NM_014673.1 /DEF=Homo sapiens KIAA0103 gene product (KIAA0103), mRNA. /FEA=mRNA /GEN=KIAA0103 /PROD=KIAA0103 gene product /DB_XREF=gi:7661909 /UG=Hs.154387 KIAA0103 gene product /FL=gb:D14659.1 gb:NM_014673.1	NM_014673		NP_055488
203585_at	0.010205	gb:NM_007150.1 /DEF=Homo sapiens zinc finger protein 185 (LIM domain) (ZNF185), mRNA. /FEA=mRNA /GEN=ZNF185 /PROD=zinc finger protein 185 (LIM domain) /DB_XREF=gi:6005971 /UG=Hs.16622 zinc finger protein 185 (LIM domain) /FL=gb:NM_007150.1	NM_007150		NP_009081
203590_at	0.018693	gb:NM_006141.1 /DEF=Homo sapiens dynein, cytoplasmic, light intermediate polypeptide 2 (DNCL12), mRNA. /FEA=mRNA /GEN=DNCL12 /PROD=dynein, cytoplasmic, light intermediate polypeptide 2 /DB_XREF=gi:5453633 /UG=Hs.194625 dynein, cytoplasmic, light intermediate polypeptide 2 /FL=gb:AF035812.1 gb:NM_006141.1	NM_006141		NP_006132
203592_s_at	0.011725	gb:NM_005860.1 /DEF=Homo sapiens follistatin-like 3 (secreted glycoprotein) (FSTL3), mRNA. /FEA=mRNA /GEN=FSTL3 /PROD=follistatin-like 3 glycoprotein /DB_XREF=gi:5031700 /UG=Hs.25348 follistatin-like 3 (secreted glycoprotein) /FL=gb:U76702.1 gb:NM_005860.1	NM_005860		NP_005851

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203594_at	7.58E-04	gb:NM_003729.1 /DEF=Homo sapiens RNA 3-terminal phosphate cyclase (RPC), mRNA. /FEA=mRNA /GEN=RPC /PROD=RNA 3-terminal phosphate cyclase /DB_XREF=gi:4506588 /UG=Hs.27076 RNA 3-terminal phosphate cyclase /FL=gb:NM_003729.1	NM_003729		NP_003720
203598_s_at	1.28E-04	Consensus includes gb:AK000979.1 /DEF=Homo sapiens cDNA FLJ10117 fis, clone HEMBA1002810, highly similar to Homo sapiens formin binding protein 21 mRNA. /FEA=mRNA /DB_XREF=gi:7021977 /UG=Hs.28307 WW domain binding protein 4 (formin binding protein 21) /FL=gb:AF071185.1 gb:NM_007187.2	NM_007187		NP_009118
203600_s_at	0.033381	gb:NM_003704.1 /DEF=Homo sapiens gene with multiple splice variants near HD locus on 4p16.3 (RES4-22), mRNA. /FEA=mRNA /GEN=RES4-22 /PROD=gene with multiple splice variants near HD locus on 4p16.3 /DB_XREF=gi:4506480 /UG=Hs.325987 gene with multiple splice variants near HD locus on 4p16.3 /FL=gb:AB000459.1 gb:NM_003704.1	NM_003704		NP_003695
203604_at	7.58E-04	KIAA0222 gene product	N38750	Hs.48450	NP_055458
203605_at	3.26E-04	gb:NM_003136.1 /DEF=Homo sapiens signal recognition particle 54kD (SRP54), mRNA. /FEA=mRNA /GEN=SRP54 /PROD=signal recognition particle 54kD /DB_XREF=gi:4507214 /UG=Hs.49346 signal recognition particle 54kD /FL=gb:U51920.1 gb:BC000652.1 gb:BC003389.1 gb:NM_003136.1	NM_003136		NP_003127

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203613_s_at	1.26E-04	gb:NM_002493.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6 (17kD, B17) (NDUFB6), mRNA. /FEA=mRNA /GEN=NDUFB6 /PROD=NADH dehydrogenase (ubiquinone) 1 betasubcomplex, 6 (17kD, B17) /DB_XREF=gi:4505364 /UG=Hs.109646 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6 (17kD, B17) /FL=gb:AF035840.1 gb:NM_002493.1 gb:AF067167.1	NM_002493		NP_002484
203615_x_at	0.026891	gb:NM_001055.1 /DEF=Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, member 1 (SULT1A1), mRNA. /FEA=mRNA /GEN=SULT1A1 /PROD=sulfotransferase family, cytosolic, 1A,phenol-preferring, member 1 /DB_XREF=gi:4507300 /UG=Hs.142 sulfotransferase family, cytosolic, 1A, phenol-preferring, member 1 /FL=gb:BC000923.2 gb:L10819.1 gb:L19955.1 gb:L19999.1 gb:NM_001055.1 gb:U09031.1 gb:U26309.1	NM_001055		NP_803880
203620_s_at	5.03E-04	gb:NM_014824.1 /DEF=Homo sapiens KIAA0769 gene product (KIAA0769), mRNA. /FEA=mRNA /GEN=KIAA0769 /PROD=KIAA0769 gene product /DB_XREF=gi:7662295 /UG=Hs.19056 KIAA0769 gene product /FL=gb:AB018312.1 gb:NM_014824.1	NM_014824		NP_055639
203630_s_at	0.001116	gb:NM_006348.1 /DEF=Homo sapiens golgi transport complex 1 (90 kDa subunit) (GOLTC1), mRNA. /FEA=mRNA /GEN=GOLTC1 /PROD=golgi transport complex 1 (90 kDa subunit) /DB_XREF=gi:5453669 /UG=Hs.239631 golgi transport complex 1 (90 kDa subunit) /FL=gb:AF058718.1 gb:NM_006348.1	NM_006348		NP_006339

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203635_at	1.26E-04	gb:NM_006052.1 /DEF=Homo sapiens Down syndrome critical region gene 3 (DSCR3), mRNA. /FEA=mRNA /GEN=DSCR3 /PROD=Down syndrome critical region protein 3 /DB_XREF=gi:5174424 /UG=Hs.26146 Down syndrome critical region gene 3 /FL=gb:D87343.1 gb:NM_006052.1	NM_006052		NP_006043
203642_s_at	0.03018	gb:NM_014900.1 /DEF=Homo sapiens KIAA0977 protein (KIAA0977), mRNA. /FEA=mRNA /GEN=KIAA0977 /PROD=KIAA0977 protein /DB_XREF=gi:7662427 /UG=Hs.300855 KIAA0977 protein /FL=gb:AB023194.1 gb:NM_014900.1	NM_014900		NP_055715
203643_at	0.003664	gb:NM_006494.1 /DEF=Homo sapiens Ets2 repressor factor (ERF), mRNA. /FEA=mRNA /GEN=ERF /PROD=Ets2 repressor factor /DB_XREF=gi:5729813 /UG=Hs.333069 Ets2 repressor factor /FL=gb:U15655.1 gb:NM_006494.1	NM_006494		NP_006485
203644_s_at	0.019292	KIAA0872 protein	AI629033	Hs.40109	NP_055755
203645_s_at	0.005852	gb:NM_004244.1 /DEF=Homo sapiens CD163 antigen (CD163), mRNA. /FEA=mRNA /GEN=CD163 /PROD=CD163 antigen /DB_XREF=gi:4758721 /UG=Hs.74076 CD163 antigen /FL=gb:NM_004244.1	NM_004244		NP_004235
203655_at	3.26E-04	gb:NM_006297.1 /DEF=Homo sapiens X-ray repair complementing defective repair in Chinese hamster cells 1 (XRCC1), mRNA. /FEA=mRNA /GEN=XRCC1 /PROD=X-ray repair cross complementing protein 1 /DB_XREF=gi:5454171 /UG=Hs.98493 X-ray repair complementing defective repair in Chinese hamster cells 1 /FL=gb:M36089.1 gb:NM_006297.1	NM_006297		NP_006288

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203656_at	3.26E-04	gb:NM_014845.1 /DEF=Homo sapiens KIAA0274 gene product (KIAA0274), mRNA. /FEA=mRNA /GEN=KIAA0274 /PROD=KIAA0274 gene product /DB_XREF=gi:7662033 /UG=Hs.10037 KIAA0274 gene product /FL=gb:D87464.1 gb:NM_014845.1	NM_014845		NP_055660
203658_at	0.003171	gb:BC001689.1 /DEF=Homo sapiens, carnitineacylcarnitine translocase, clone MGC:1207, mRNA, complete cds. /FEA=mRNA /PROD=carnitineacylcarnitine translocase /DB_XREF=gi:12804552 /UG=Hs.13845 solute carrier family 25 (carnitineacylcarnitine translocase), member 20 /FL=gb:BC001689.1 gb:NM_000387.2	BC001689		NP_000378
203661_s_at	0.002536	gb:BC002660.1 /DEF=Homo sapiens, tropomodulin, clone MGC:3643, mRNA, complete cds. /FEA=mRNA /PROD=tropomodulin /DB_XREF=gi:12803650 /UG=Hs.170453 tropomodulin /FL=gb:BC002660.1 gb:M77016.1 gb:NM_003275.1	BC002660		NP_003266
203663_s_at	0.005852	gb:NM_004255.1 /DEF=Homo sapiens cytochrome c oxidase subunit Va (COX5A), mRNA. /FEA=mRNA /GEN=COX5A /PROD=cytochrome c oxidase subunit Va /DB_XREF=gi:4758037 /UG=Hs.181028 cytochrome c oxidase subunit Va /FL=gb:NM_004255.1 gb:M22760.1	NM_004255		NP_004246

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203665_at	0.00228	gb:NM_002133.1 /DEF=Homo sapiens heme oxygenase (decycling) 1 (HMOX1), mRNA. /FEA=mRNA /GEN=HMOX1 /PROD=heme oxygenase (decycling) 1 /DB_XREF=gi:4504436 /UG=Hs.202833 heme oxygenase (decycling) 1 /FL=gb:NM_002133.1	NM_002133		NP_002124
203667_at	0.007779	gb:NM_004607.1 /DEF=Homo sapiens tubulin-specific chaperone a (TBCA), mRNA. /FEA=mRNA /GEN=TBCA /PROD=beta-tubulin cofactor A /DB_XREF=gi:4759211 /UG=Hs.24930 tubulin-specific chaperone a /FL=gb:AF038952.1 gb:NM_004607.1	NM_004607		NP_004598
203668_at	0.016934	gb:NM_006715.1 /DEF=Homo sapiens mannosidase, alpha, class 2C, member 1 (MAN2C1), mRNA. /FEA=mRNA /GEN=MAN2C1 /PROD=mannosidase, alpha 6A8 /DB_XREF=gi:6631092 /UG=Hs.26232 mannosidase, alpha, class 2C, member 1 /FL=gb:U37248.1 gb:AF044414.2 gb:NM_006715.1	NM_006715		NP_006706
203669_s_at	0.007052	gb:NM_012079.2 /DEF=Homo sapiens diacylglycerol O-acyltransferase (mouse) homolog (DGAT), mRNA. /FEA=mRNA /GEN=DGAT /PROD=diacylglycerol O-acyltransferase (mouse)homolog /DB_XREF=gi:7382489 /UG=Hs.288627 diacylglycerol O-acyltransferase (mouse) homolog /FL=gb:AF059202.1 gb:NM_012079.2	NM_012079		NP_036211

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203672_x_at	0.004075	gb:U12387.1 /DEF=Human thiopurine methyltransferase (TPMT) mRNA, complete cds. /FEA=mRNA /GEN=TPMT /PROD=thiopurine methyltransferase /DB_XREF=gi:805083 /UG=Hs.296922 thiopurine S-methyltransferase /FL=gb:BC005339.1 gb:NM_000367.1 gb:U12387.1	U12387		NP_000358
203675_at	1.26E-04	gb:NM_005013.1 /DEF=Homo sapiens nucleobindin 2 (NUCB2), mRNA. /FEA=mRNA /GEN=NUCB2 /PROD=nucleobindin 2 /DB_XREF=gi:4826869 /UG=Hs.3164 nucleobindin 2 /FL=gb:AF052642.1 gb:AF052643.1 gb:AF052644.1 gb:NM_005013.1	NM_005013		NP_005004
203685_at	0.041055	gb:NM_000633.1 /DEF=Homo sapiens B-cell CLLlymphoma 2 (BCL2), nuclear gene encoding mitochondrial protein, transcript variant alpha, mRNA. /FEA=mRNA /GEN=BCL2 /PROD=B-cell lymphoma protein 2 alpha /DB_XREF=gi:4557354 /UG=Hs.79241 B-cell CLLlymphoma 2 /FL=gb:M13994.1 gb:NM_000633.1	NM_000633		NP_000648
203690_at	2.05E-04	gb:NM_006322.1 /DEF=Homo sapiens spindle pole body protein (GCP3), mRNA. /FEA=mRNA /GEN=GCP3 /PROD=spindle pole body protein /DB_XREF=gi:5453659 /UG=Hs.9884 spindle pole body protein /FL=gb:AF042378.1 gb:NM_006322.1	NM_006322		NP_006313

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203691_at	0.003171	gb:NM_002638.1 /DEF=Homo sapiens protease inhibitor 3, skin-derived (SKALP) (PI3), mRNA. /FEA=mRNA /GEN=PI3 /PROD=protease inhibitor 3, skin-derived (SKALP) /DB_XREF=gi:4505786 /UG=Hs.112341 protease inhibitor 3, skin-derived (SKALP) /FL=gb:NM_002638.1	NM_002638		NP_002629
203693_s_at	0.00228	gb:NM_001949.2 /DEF=Homo sapiens E2F transcription factor 3 (E2F3) mRNA, complete cds. /FEA=mRNA /GEN=E2F3 /PROD=E2F transcription factor 3 /DB_XREF=gi:12669913 /UG=Hs.1189 E2F transcription factor 3 /FL=gb:NM_001949.2	NM_001949		NP_001940
203704_s_at	3.26E-04	ras responsive element binding protein 1	AW118862	Hs.171942	NP_002946
203707_at	0.013223	gb:NM_005741.1 /DEF=Homo sapiens zinc finger protein 263 (ZNF263), mRNA. /FEA=mRNA /GEN=ZNF263 /PROD=zinc finger protein 263 /DB_XREF=gi:5032240 /UG=Hs.182528 zinc finger protein 263 /FL=gb:D88827.1 gb:NM_005741.1	NM_005741		NP_005732
203713_s_at	6.68E-04	gb:NM_004524.1 /DEF=Homo sapiens lethal giant larvae (Drosophila) homolog 2 (LLGL2), mRNA. /FEA=mRNA /GEN=LLGL2 /PROD=lethal giant larvae (Drosophila) homolog 2 /DB_XREF=gi:4758679 /UG=Hs.3123 lethal giant larvae (Drosophila) homolog 2 /FL=gb:NM_004524.1	NM_004524		NP_004515
203721_s_at	0.00228	gb:NM_016001.1 /DEF=Homo sapiens CGI-48 protein (LOC51096), mRNA. /FEA=mRNA /GEN=LOC51096 /PROD=CGI-48 protein /DB_XREF=gi:7705764 /UG=Hs.6153 CGI-48 protein /FL=gb:AF151806.1 gb:NM_016001.1	NM_016001		NP_057085

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203723_at	0.02145	gb:NM_002221.1 /DEF=Homo sapiens inositol 1,4,5-trisphosphate 3-kinase B (ITPKB), mRNA. /FEA=mRNA /GEN=ITPKB /PROD=1D-myo-inositol-trisphosphate 3-kinase B /DB_XREF=gi:4504790 /UG=Hs.78877 inositol 1,4,5-trisphosphate 3-kinase B /FL=gb:NM_002221.1	NM_002221		NP_002212
203725_at	1.26E-04	gb:NM_001924.2 /DEF=Homo sapiens growth arrest and DNA-damage-inducible, alpha (GADD45A), mRNA. /FEA=mRNA /GEN=GADD45A /PROD=growth arrest and DNA-damage-inducible, alpha /DB_XREF=gi:9790904 /UG=Hs.80409 growth arrest and DNA-damage-inducible, alpha /FL=gb:M60974.1 gb:NM_001924.2	NM_001924		NP_001915
203730_s_at	0.046687	zinc finger protein 95 homolog (mouse)	BF196931	Hs.110839	NP_659570
203731_s_at	0.048086	gb:NM_014569.1 /DEF=Homo sapiens zinc finger protein homologous to Zfp95 in mouse (ZFP95), mRNA. /FEA=mRNA /GEN=ZFP95 /PROD=zinc finger protein homologous to Zfp95 inmouse /DB_XREF=gi:11036641 /UG=Hs.110839 zinc finger protein homologous to Zfp95 in mouse /FL=gb:NM_014569.1 gb:AB023232.1	NM_014569		NP_659570
203732_at	0.001116	gb:NM_016213.1 /DEF=Homo sapiens thyroid hormone receptor interactor 4 (TRIP4), mRNA. /FEA=mRNA /GEN=TRIP4 /PROD=activating signal cointegrator 1 /DB_XREF=gi:7706430 /UG=Hs.116784 thyroid hormone receptor interactor 4 /FL=gb:AF168418.1 gb:NM_016213.1	NM_016213		NP_057297
203738_at	0.033381	hypothetical protein FLJ11193	AI421192	Hs.151046	NP_060826

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203739_at	2.05E-04	gb:NM_006526.1 /DEF=Homo sapiens zinc finger protein 217 (ZNF217), mRNA. /FEA=mRNA /GEN=ZNF217 /PROD=zinc finger protein 217 /DB_XREF=gi:5730123 /UG=Hs.155040 zinc finger protein 217 /FL=gb:AF041259.1 gb:NM_006526.1	NM_006526		NP_006517
203742_s_at	0.022795	thymine-DNA glycosylase	BF674842	Hs.173824	
203743_s_at	0.007779	gb:NM_003211.1 /DEF=Homo sapiens thymine-DNA glycosylase (TDG), mRNA. /FEA=mRNA /GEN=TDG /PROD=thymine-DNA glycosylase /DB_XREF=gi:4507422 /UG=Hs.173824 thymine-DNA glycosylase /FL=gb:U51166.1 gb:NM_003211.1	NM_003211		NP_003202
203744_at	3.56E-04	gb:NM_005342.1 /DEF=Homo sapiens high-mobility group (nonhistone chromosomal) protein 4 (HMG4), mRNA. /FEA=mRNA /GEN=HMG4 /PROD=high-mobility group (nonhistone chromosomal)protein 4 /DB_XREF=gi:4885420 /UG=Hs.19114 high-mobility group (nonhistone chromosomal) protein 4 /FL=gb:NM_005342.1	NM_005342		NP_005333
203746_s_at	0.00136	gb:NM_005333.1 /DEF=Homo sapiens holocytochrome c synthase (cytochrome c heme-lyase) (HCCS), mRNA. /FEA=mRNA /GEN=HCCS /PROD=holocytochrome c synthase (cytochrome cheme-lyase) /DB_XREF=gi:4885400 /UG=Hs.211571 holocytochrome c synthase (cytochrome c heme-lyase) /FL=gb:U36787.1 gb:NM_005333.1	NM_005333		NP_005324

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203748_x_at	2.05E-04	gb:NM_016839.1 /DEF=Homo sapiens RNA binding motif, single stranded interacting protein 1 (RBMS1), transcript variant MSSP-2, mRNA. /FEA=mRNA /GEN=RBMS1 /PROD=RNA binding motif, single stranded interactingprotein 1, isoform b /DB_XREF=gi:8400723 /UG=Hs.241567 RNA binding motif, single stranded interacting protein 1 /FL=gb:NM_016839.1	NM_016839		NP_058523
203757_s_at	0.041795	gb:BC005008.1 /DEF=Homo sapiens, carcinoembryonic antigen related cell adhesion molecule 6 (non-specific cross reacting antigen), clone MGC:10467, mRNA, complete cds. /FEA=mRNA /PROD=carcinoembryonic antigen-related cell adhesionmolecule 6 (non-specific cross reacting antigen) /DB_XREF=gi:13477106 /UG=Hs.73848 carcinoembryonic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen) /FL=gb:BC005008.1 gb:M18216.1 gb:M29541.1 gb:NM_002483.1	BC005008		NP_002474
203761_at	0.00228	gb:NM_006748.1 /DEF=Homo sapiens Src-like-adapter (SLA), mRNA. /FEA=mRNA /GEN=SLA /PROD=Src-like-adapter /DB_XREF=gi:5803170 /UG=Hs.75367 Src-like-adapter /FL=gb:U30473.1 gb:D89077.1 gb:U44403.1 gb:NM_006748.1	NM_006748		NP_006739
203764_at	0.002536	gb:NM_014750.1 /DEF=Homo sapiens KIAA0008 gene product (KIAA0008), mRNA. /FEA=mRNA /GEN=KIAA0008 /PROD=KIAA0008 gene product /DB_XREF=gi:7661851 /UG=Hs.77695 KIAA0008 gene product /FL=gb:D13633.1 gb:NM_014750.1	NM_014750		NP_055565

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203765_at	0.010205	gb:NM_012198.1 /DEF=Homo sapiens grancalcin (GCL), mRNA. /FEA=mRNA /GEN=GCL /PROD=grancalcin /DB_XREF=gi:6912387 /UG=Hs.79381 grancalcin /FL=gb:BC005214.1 gb:M81637.1 gb:NM_012198.1	NM_012198		NP_036330
203773_x_at	0.013223	gb:NM_000712.1 /DEF=Homo sapiens biliverdin reductase A (BLVRA), mRNA. /FEA=mRNA /GEN=BLVRA /PROD=biliverdin reductase A /DB_XREF=gi:4502416 /UG=Hs.81029 biliverdin reductase A /FL=gb:U34877.1 gb:NM_000712.1	NM_000712		NP_000703
203775_at	3.26E-04	gb:NM_014251.1 /DEF=Homo sapiens solute carrier family 25, member 13 (citrin) (SLC25A13), mRNA. /FEA=mRNA /GEN=SLC25A13 /PROD=solute carrier family 25, member 13 (citrin) /DB_XREF=gi:7657580 /UG=Hs.9599 solute carrier family 25, member 13 (citrin) /FL=gb:AF118838.1 gb:NM_014251.1	NM_014251		NP_055066
203776_at	0.033381	gb:NM_015698.1 /DEF=Homo sapiens T54 protein (T54), mRNA. /FEA=mRNA /GEN=T54 /PROD=T54 protein /DB_XREF=gi:7662668 /UG=Hs.100391 T54 protein /FL=gb:BC000397.1 gb:BC003148.1 gb:U66359.1 gb:NM_015698.1	NM_015698		NP_056513
203778_at	5.03E-04	gb:NM_005908.1 /DEF=Homo sapiens mannosidase, beta A, lysosomal (MANBA), mRNA. /FEA=mRNA /GEN=MANBA /PROD=mannosidase, beta A, lysosomal /DB_XREF=gi:5174522 /UG=Hs.115945 mannosidase, beta A, lysosomal /FL=gb:U60337.1 gb:NM_005908.1	NM_005908		NP_005899

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203781_at	0.003171	gb:NM_004891.1 /DEF=Homo sapiens chromosome 2 open reading frame 1 (C2ORF1), mRNA. /FEA=mRNA /GEN=C2ORF1 /PROD=chromosome 2 open reading frame 1 /DB_XREF=gi:4759047 /UG=Hs.14454 chromosome 2 open reading frame 1 /FL=gb:AF047440.1 gb:NM_004891.1	NM_004891		NP_663303
203789_s_at	0.035611	gb:NM_006379.1 /DEF=Homo sapiens sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C (SEMA3C), mRNA. /FEA=mRNA /GEN=SEMA3C /PROD=sema domain, immunoglobulin domain (Ig), shortbasic domain, secreted, (semaphorin) 3C /DB_XREF=gi:5454047 /UG=Hs.171921 sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C /FL=gb:AB000220.1 gb:NM_006379.1	NM_006379		NP_006370
203799_at	0.005852	gb:NM_014880.1 /DEF=Homo sapiens KIAA0022 gene product (KIAA0022), mRNA. /FEA=mRNA /GEN=KIAA0022 /PROD=KIAA0022 gene product /DB_XREF=gi:7661867 /UG=Hs.2441 KIAA0022 gene product /FL=gb:D14664.1 gb:NM_014880.1	NM_014880		NP_055695
203800_s_at	5.03E-04	Consensus includes gb:BG254653 /FEA=EST /DB_XREF=gi:12764469 /DB_XREF=est:602368621F1 /CLONE=IMAGE:4476773 /UG=Hs.247324 hypothetical protein LOC63931 /FL=gb:NM_022100.1	NM_022100		NP_071383
203810_at	0.034221	DnaJ (Hsp40) homolog, subfamily B, member 4	BG252490	Hs.41693	NP_008965

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203814_s_at	0.00161	gb:NM_000904.1 /DEF=Homo sapiens NAD(P)H menadione oxidoreductase 2, dioxin-inducible (NMOR2), mRNA. /FEA=mRNA /GEN=NMOR2 /PROD=NAD(P)H menadione oxidoreductase 2,dioxin-inducible /DB_XREF=gi:4505416 /UG=Hs.73956 NAD(P)H menadione oxidoreductase 2, dioxin-inducible /FL=gb:J02888.1 gb:NM_000904.1	NM_000904		NP_000895
203825_at	0.002418	gb:NM_007371.2 /DEF=Homo sapiens bromodomain-containing 3 (BRD3), mRNA. /FEA=mRNA /GEN=BRD3 /PROD=bromodomain-containing protein 3 /DB_XREF=gi:12408642 /UG=Hs.86896 bromodomain-containing 3 /FL=gb:NM_007371.2 gb:D26362.1	NM_007371		NP_031397
203827_at	5.03E-04	gb:NM_017983.1 /DEF=Homo sapiens hypothetical protein FLJ10055 (FLJ10055), mRNA. /FEA=mRNA /GEN=FLJ10055 /PROD=hypothetical protein FLJ10055 /DB_XREF=gi:8922207 /UG=Hs.9398 hypothetical protein FLJ10055 /FL=gb:NM_017983.1	NM_017983		NP_060453
203837_at	0.003171	gb:NM_005923.2 /DEF=Homo sapiens mitogen-activated protein kinase kinase kinase 5 (MAP3K5), mRNA. /FEA=mRNA /GEN=MAP3K5 /PROD=MAPERK kinase kinase 5 /DB_XREF=gi:6031181 /UG=Hs.151988 mitogen-activated protein kinase kinase kinase 5 /FL=gb:U67156.1 gb:D84476.1 gb:NM_005923.2	NM_005923		NP_005914

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203855_at	0.041055	gb:NM_014969.1 /DEF=Homo sapiens KIAA0893 protein (KIAA0893), mRNA. /FEA=mRNA /GEN=KIAA0893 /PROD=KIAA0893 protein /DB_XREF=gi:7662363 /UG=Hs.3830 KIAA0893 protein /FL=gb:AB020700.1 gb:NM_014969.1	NM_014969		NP_055784
203856_at	0.011788	gb:NM_003384.1 /DEF=Homo sapiens vaccinia related kinase 1 (VRK1), mRNA. /FEA=mRNA /GEN=VRK1 /PROD=vaccinia related kinase 1 /DB_XREF=gi:4507902 /UG=Hs.48269 vaccinia related kinase 1 /FL=gb:AB000449.1 gb:NM_003384.1	NM_003384		NP_003375
203857_s_at	0.001303	gb:NM_006810.1 /DEF=Homo sapiens for protein disulfide isomerase-related (PDIR), mRNA. /FEA=mRNA /GEN=PDIR /PROD=for protein disulfide isomerase-related /DB_XREF=gi:5803120 /UG=Hs.76901 for protein disulfide isomerase-related /FL=gb:D49490.1 gb:NM_006810.1	NM_006810		NP_006801
203880_at	0.001116	gb:NM_005694.1 /DEF=Homo sapiens COX17 (yeast) homolog, cytochrome c oxidase assembly protein (COX17), mRNA. /FEA=mRNA /GEN=COX17 /PROD=COX17 (yeast) homolog, cytochrome c oxidase assembly protein /DB_XREF=gi:5031644 /UG=Hs.16297 COX17 (yeast) homolog, cytochrome c oxidase assembly protein /FL=gb:L77701.1 gb:NM_005694.1	NM_005694		NP_005685

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203884_s_at	0.00161	gb:NM_014904.1 /DEF=Homo sapiens KIAA0941 protein (KIAA0941), mRNA. /FEA=mRNA /GEN=KIAA0941 /PROD=KIAA0941 protein /DB_XREF=gi:7662393 /UG=Hs.173656 KIAA0941 protein /FL=gb:AB023158.1 gb:NM_014904.1	NM_014904		NP_055719
203885_at	1.26E-04	gb:NM_014999.1 /DEF=Homo sapiens KIAA0118 protein (KIAA0118), mRNA. /FEA=mRNA /GEN=KIAA0118 /PROD=KIAA0118 protein /DB_XREF=gi:7661921 /UG=Hs.184627 KIAA0118 protein /FL=gb:AF091035.1 gb:NM_014999.1	NM_014999		NP_055814
203887_s_at	7.58E-04	gb:NM_000361.1 /DEF=Homo sapiens thrombomodulin (THBD), mRNA. /FEA=mRNA /GEN=THBD /PROD=thrombomodulin /DB_XREF=gi:4507482 /UG=Hs.2030 thrombomodulin /FL=gb:M16552.1 gb:NM_000361.1	NM_000361		NP_000352
203888_at	5.96E-04	gb:NM_000361.1 /DEF=Homo sapiens thrombomodulin (THBD), mRNA. /FEA=mRNA /GEN=THBD /PROD=thrombomodulin /DB_XREF=gi:4507482 /UG=Hs.2030 thrombomodulin /FL=gb:M16552.1 gb:NM_000361.1	NM_000361		NP_000352
203897_at	7.58E-04	hypothetical protein A-211C6.1	BE963444	Hs.28607	NP_065157

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203904_x_at	0.003942	gb:NM_002231.2 /DEF=Homo sapiens kangai 1 (suppression of tumorigenicity 6, prostate; CD82 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4)) (KAI1), mRNA. /FEA=mRNA /GEN=KAI1 /PROD=kangai 1 /DB_XREF=gi:13259537 /UG=Hs.323949 kangai 1 (suppression of tumorigenicity 6, prostate; CD82 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4)) /FL=gb:BC000726.1 gb:BC001821.1 gb:NM_002231.2 gb:U20770.1	NM_002231		NP_002222
203906_at	1.26E-04	KIAA0763 gene product	AI652645	Hs.409124	NP_055684
203907_s_at	1.26E-04	gb:NM_014869.1 /DEF=Homo sapiens KIAA0763 gene product (KIAA0763), mRNA. /FEA=mRNA /GEN=KIAA0763 /PROD=KIAA0763 gene product /DB_XREF=gi:7662289 /UG=Hs.4764 KIAA0763 gene product /FL=gb:AB018306.1 gb:NM_014869.1	NM_014869		NP_055684
203909_at	1.26E-04	gb:NM_006359.1 /DEF=Homo sapiens solute carrier family 9 (sodiumhydrogen exchanger), isoform 6 (SLC9A6), mRNA. /FEA=mRNA /GEN=SLC9A6 /PROD=solute carrier family 9 (sodiumhydrogenexchanger), isoform 6 /DB_XREF=gi:5454069 /UG=Hs.62185 solute carrier family 9 (sodiumhydrogen exchanger), isoform 6 /FL=gb:AF030409.1 gb:NM_006359.1	NM_006359		NP_006350

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203912_s_at	1.36E-04	gb:NM_006730.1 /DEF=Homo sapiens deoxyribonuclease I-like 1 (DNASE1L1), mRNA. /FEA=mRNA /GEN=DNASE1L1 /PROD=deoxyribonuclease I-like 1 /DB_XREF=gi:5803006 /UG=Hs.77091 deoxyribonuclease I-like 1 /FL=gb:U06846.1 gb:NM_006730.1	NM_006730		NP_006721
203914_x_at	0.021448	gb:NM_000860.1 /DEF=Homo sapiens hydroxyprostaglandin dehydrogenase 15-(NAD) (HPGD), mRNA. /FEA=mRNA /GEN=HPGD /PROD=hydroxyprostaglandin dehydrogenase 15-(NAD) /DB_XREF=gi:4504478 /UG=Hs.77348 hydroxyprostaglandin dehydrogenase 15-(NAD) /FL=gb:L76465.1 gb:NM_000860.1	NM_000860		NP_000851
203922_s_at	0.016934	cytochrome b-245, beta polypeptide (chronic granulomatous disease)	AI308863	Hs.88974	NP_000388
203925_at	1.26E-04	gb:NM_002061.1 /DEF=Homo sapiens glutamate-cysteine ligase, modifier subunit (GCLM), mRNA. /FEA=mRNA /GEN=GCLM /PROD=glutamate-cysteine ligase regulatory protein /DB_XREF=gi:4504010 /UG=Hs.89709 glutamate-cysteine ligase, modifier subunit /FL=gb:NM_002061.1 gb:L35546.1	NM_002061		NP_002052
203932_at	0.016934	gb:NM_002118.1 /DEF=Homo sapiens major histocompatibility complex, class II, DM beta (HLA-DMB), mRNA. /FEA=mRNA /GEN=HLA-DMB /PROD=major histocompatibility complex, class II, DMbeta /DB_XREF=gi:4504398 /UG=Hs.1162 major histocompatibility complex, class II, DM beta /FL=gb:NM_002118.1 gb:U15085.1	NM_002118		NP_002109

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203936_s_at	0.00228	gb:NM_004994.1 /DEF=Homo sapiens matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase) (MMP9), mRNA. /FEA=mRNA /GEN=MMP9 /PROD=matrix metalloproteinase 9 preproprotein /DB_XREF=gi:4826835 /UG=Hs.151738 matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase) /FL=gb:J05070.1 gb:NM_004994.1	NM_004994		NP_004985
203943_at	0.033381	gb:NM_004798.1 /DEF=Homo sapiens kinesin family member 3B (KIF3B), mRNA. /FEA=mRNA /GEN=KIF3B /PROD=kinesin family member 3B /DB_XREF=gi:4758645 /UG=Hs.168212 kinesin family member 3B /FL=gb:AB002357.1 gb:NM_004798.1	NM_004798		NP_004789
203945_at	0.00136	gb:NM_001172.2 /DEF=Homo sapiens arginase, type II (ARG2), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=ARG2 /PROD=arginase, type II precursor /DB_XREF=gi:10947110 /UG=Hs.172851 arginase, type II /FL=gb:NM_001172.2 gb:BC001350.1 gb:D86724.1 gb:U75667.1 gb:U82256.1	NM_001172		NP_001163
203949_at	0.011725	gb:NM_000250.1 /DEF=Homo sapiens myeloperoxidase (MPO), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=MPO /PROD=myeloperoxidase /DB_XREF=gi:4557758 /UG=Hs.1817 myeloperoxidase /FL=gb:M19507.1 gb:J02694.1 gb:NM_000250.1	NM_000250		NP_000241
203958_s_at	0.009689	KIAA0478 gene product	AI557467	Hs.4236	NP_055685

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203960_s_at	0.006096	gb:NM_016126.1 /DEF=Homo sapiens HSPCO34 protein (LOC51668), mRNA. /FEA=mRNA /GEN=LOC51668 /PROD=HSPCO34 protein /DB_XREF=gi:7706382 /UG=Hs.46967 HSPCO34 protein /FL=gb:BC005245.1 gb:AF100747.1 gb:NM_016126.1	NM_016126		NP_057210
203962_s_at	0.023856	gb:NM_006393.1 /DEF=Homo sapiens nebulette (NEBL), mRNA. /FEA=mRNA /GEN=NEBL /PROD=nebulette /DB_XREF=gi:5453757 /UG=Hs.5025 nebulette /FL=gb:NM_006393.1	NM_006393		NP_006384
203966_s_at	1.26E-04	gb:NM_021003.1 /DEF=Homo sapiens protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform (PPM1A), mRNA. /FEA=mRNA /GEN=PPM1A /PROD=protein phosphatase 1A (formerly 2C),magnesium-dependent, alpha isoform /DB_XREF=gi:10337594 /UG=Hs.57764 protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform /FL=gb:NM_021003.1	NM_021003		NP_808821
203968_s_at	0.029713	gb:NM_001254.1 /DEF=Homo sapiens CDC6 (cell division cycle 6, S. cerevisiae) homolog (CDC6), mRNA. /FEA=mRNA /GEN=CDC6 /PROD=CDC6 (cell division cycle 6, S. cerevisiae)homolog /DB_XREF=gi:4502702 /UG=Hs.69563 CDC6 (cell division cycle 6, S. cerevisiae) homolog /FL=gb:U77949.1 gb:AF022109.1 gb:NM_001254.1	NM_001254		NP_001245

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203971_at	0.011788	gb:NM_001859.1 /DEF=Homo sapiens solute carrier family 31 (copper transporters), member 1 (SLC31A1), mRNA. /FEA=mRNA /GEN=SLC31A1 /PROD=solute carrier family 31 (copper transporters),member 1 /DB_XREF=gi:4507014 /UG=Hs.73614 solute carrier family 31 (copper transporters), member 1 /FL=gb:U83460.1 gb:NM_001859.1	NM_001859		NP_001850
203974_at	0.007779	gb:NM_012080.1 /DEF=Homo sapiens DNA segment, numerous copies, expressed probes (GS1 gene) (DXF68S1E), mRNA. /FEA=mRNA /GEN=DXF68S1E /PROD=DNA segment, numerous copies, expressed probes(GS1 gene) /DB_XREF=gi:6912345 /UG=Hs.78991 DNA segment, numerous copies, expressed probes (GS1 gene) /FL=gb:M86934.1 gb:NM_012080.1	NM_012080		NP_036212
203983_at	0.003171	gb:NM_005999.1 /DEF=Homo sapiens translin-associated factor X (TSNAX), mRNA. /FEA=mRNA /GEN=TSNAX /PROD=translin-associated factor X /DB_XREF=gi:5174730 /UG=Hs.96247 translin-associated factor X /FL=gb:NM_005999.1	NM_005999		NP_005990
203984_s_at	0.013223	gb:U60521.1 /DEF=Human protease proMch6 (Mch6) mRNA, complete cds. /FEA=mRNA /GEN=Mch6 /PROD=proMch6 /DB_XREF=gi:1532150 /UG=Hs.100641 caspase 9, apoptosis-related cysteine protease /FL=gb:BC002452.1 gb:U56390.1 gb:U60521.1 gb:NM_001229.1	U60521		NP_127463

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
203985_at	0.013088	gb:NM_012256.1 /DEF=Homo sapiens zinc finger protein 212 (ZNF212), mRNA. /FEA=mRNA /GEN=ZNF212 /PROD=zinc finger protein 212 /DB_XREF=gi:6912749 /UG=Hs.108139 zinc finger protein 212 /FL=gb:U38864.1 gb:NM_012256.1	NM_012256		NP_036388
203992_s_at	0.00161	gb:AF000992.1 /DEF=Homo sapiens ubiquitous TPR motif, X isoform (UTX) mRNA, alternative transcript 1, complete cds. /FEA=mRNA /GEN=UTX /PROD=ubiquitous TPR motif, X isoform /DB_XREF=gi:2580569 /UG=Hs.13980 ubiquitously transcribed tetratricopeptide repeat gene, X chromosome /FL=gb:NM_021140.1 gb:AF000992.1 gb:AF000993.1	AF000992		NP_066963
204001_at	0.010205	gb:NM_003084.1 /DEF=Homo sapiens small nuclear RNA activating complex, polypeptide 3, 50kD (SNAPC3), mRNA. /FEA=mRNA /GEN=SNAPC3 /PROD=small nuclear RNA activating complex, polypeptide 3, 50kD /DB_XREF=gi:4507104 /UG=Hs.164915 small nuclear RNA activating complex, polypeptide 3, 50kD /FL=gb:U66413.1 gb:NM_003084.1	NM_003084		NP_003075
204009_s_at	1.26E-04	v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog	W80678	Hs.351221	NP_203524
204018_x_at	5.03E-04	gb:NM_000558.2 /DEF=Homo sapiens hemoglobin, alpha 1 (HBA1), mRNA. /FEA=mRNA /GEN=HBA1 /PROD=hemoglobin, alpha 1 /DB_XREF=gi:6715603 /UG=Hs.251577 hemoglobin, alpha 1 /FL=gb:NM_000558.2	NM_000558		NP_000549
204022_at	1.26E-04	Nedd-4-like ubiquitin-protein ligase	AI668780	Hs.333382	NP_008945

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204026_s_at	2.05E-04	gb:NM_007057.1 /DEF=Homo sapiens ZW10 interactor (ZWINT), mRNA. /FEA=mRNA /GEN=ZWINT /PROD=ZW10 interactor /DB_XREF=gi:6857828 /UG=Hs.42650 ZW10 interactor /FL=gb:AF067656.1 gb:NM_007057.1	NM_007057		NP_127490
204028_s_at	3.26E-04	gb:NM_012197.2 /DEF=Homo sapiens rab6 GTPase activating protein (GAP and centrosome-associated) (GAPCENA), mRNA. /FEA=mRNA /GEN=GAPCENA /PROD=rab6 GTPase activating protein (GAP and centrosome-associated) /DB_XREF=gi:12232372 /UG=Hs.55099 rab6 GTPase activating protein (GAP and centrosome-associated) /FL=gb:NM_012197.2	NM_012197		NP_036329
204031_s_at	0.00228	gb:NM_005016.1 /DEF=Homo sapiens poly(rC)-binding protein 2 (PCBP2), mRNA. /FEA=mRNA /GEN=PCBP2 /PROD=poly(rC)-binding protein 2 /DB_XREF=gi:4826885 /UG=Hs.63525 poly(rC)-binding protein 2 /FL=gb:BC001155.1 gb:NM_005016.1	NM_005016		NP_114366
204034_at	1.26E-04	gb:NM_014297.1 /DEF=Homo sapiens protein expressed in thyroid (YF13H12), mRNA. /FEA=mRNA /GEN=YF13H12 /PROD=protein expressed in thyroid /DB_XREF=gi:7657686 /UG=Hs.7486 protein expressed in thyroid /FL=gb:NM_014297.1	NM_014297		NP_055112

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204038_s_at	0.005852	gb:NM_001401.1 /DEF=Homo sapiens endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2 (EDG2), mRNA. /FEA=mRNA /GEN=EDG2 /PROD=endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2 /DB_XREF=gi:4503456 /UG=Hs.75794 endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2 /FL=gb:U78192.1 gb:U80811.1 gb:NM_001401.1	NM_001401		NP_476500
204046_at	0.04734	gb:NM_004573.1 /DEF=Homo sapiens phospholipase C, beta 2 (PLCB2), mRNA. /FEA=mRNA /GEN=PLCB2 /PROD=phospholipase C, beta 2 /DB_XREF=gi:4758937 /UG=Hs.994 phospholipase C, beta 2 /FL=gb:M95678.1 gb:NM_004573.1	NM_004573		NP_004564
204050_s_at	1.26E-04	gb:NM_001833.1 /DEF=Homo sapiens clathrin, light polypeptide (Lca) (CLTA), transcript variant nonbrain, mRNA. /FEA=mRNA /GEN=CLTA /PROD=clathrin, light polypeptide A (Lca) isoform a /DB_XREF=gi:4502898 /UG=Hs.104143 clathrin, light polypeptide (Lca) /FL=gb:M20472.1 gb:NM_001833.1	NM_001833		NP_009027
204069_at	0.025958	gb:NM_002398.1 /DEF=Homo sapiens Meis1 (mouse) homolog (MEIS1), mRNA. /FEA=mRNA /GEN=MEIS1 /PROD=Meis1 homolog /DB_XREF=gi:4505150 /UG=Hs.170177 Meis1 (mouse) homolog /FL=gb:U85707.1 gb:NM_002398.1	NM_002398		NP_002389

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204079_at	3.26E-04	gb:NM_003595.1 /DEF=Homo sapiens tyrosylprotein sulfotransferase 2 (TPST2), mRNA. /FEA=mRNA /GEN=TPST2 /PROD=tyrosylprotein sulfotransferase 2 /DB_XREF=gi:4507666 /UG=Hs.26350 tyrosylprotein sulfotransferase 2 /FL=gb:AL136623.1 gb:BC001057.1 gb:AF061254.1 gb:AF049891.1 gb:NM_003595.1	NM_003595		NP_003586
204082_at	0.001116	gb:NM_006195.1 /DEF=Homo sapiens pre-B-cell leukemia transcription factor 3 (PBX3), mRNA. /FEA=mRNA /GEN=PBX3 /PROD=pre-B-cell leukemia transcription factor 3 /DB_XREF=gi:5453851 /UG=Hs.294101 pre-B-cell leukemia transcription factor 3 /FL=gb:NM_006195.1	NM_006195		NP_006186
204084_s_at	0.033381	ceroid-lipofuscinosis, neuronal 5	AI911687	Hs.30213	NP_006484
204091_at	0.00423	gb:NM_002601.1 /DEF=Homo sapiens phosphodiesterase 6D, cGMP-specific, rod, delta (PDE6D), mRNA. /FEA=mRNA /GEN=PDE6D /PROD=phosphodiesterase 6D, cGMP-specific, rod, delta /DB_XREF=gi:4505670 /UG=Hs.48291 phosphodiesterase 6D, cGMP-specific, rod, delta /FL=gb:AF022912.1 gb:NM_002601.1	NM_002601		NP_002592
204093_at	0.013223	gb:NM_001239.1 /DEF=Homo sapiens cyclin H (CCNH), mRNA. /FEA=mRNA /GEN=CCNH /PROD=cyclin H /DB_XREF=gi:4502622 /UG=Hs.514 cyclin H /FL=gb:BC005280.1 gb:NM_001239.1 gb:U12685.1 gb:U11791.1	NM_001239		NP_001230

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204099_at	0.00228	gb:NM_003078.1 /DEF=Homo sapiens SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 3 (SMARCD3), mRNA. /FEA=mRNA /GEN=SMARCD3 /PROD=SWISNF related, matrix associated, actindependent regulator of chromatin, subfamily d, member 3 /DB_XREF=gi:4507086 /UG=Hs.71622 SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 3 /FL=gb:BC002628.1 gb:U66619.1 gb:NM_003078.1	NM_003078		NP_003069
204101_at	0.041795	gb:NM_000252.1 /DEF=Homo sapiens myotubular myopathy 1 (MTM1), mRNA. /FEA=mRNA /GEN=MTM1 /PROD=myotubularin /DB_XREF=gi:4557895 /UG=Hs.75302 myotubular myopathy 1 /FL=gb:U46024.1 gb:NM_000252.1	NM_000252		NP_000243
204116_at	0.003171	gb:NM_000206.1 /DEF=Homo sapiens interleukin 2 receptor, gamma (severe combined immunodeficiency) (IL2RG), mRNA. /FEA=mRNA /GEN=IL2RG /PROD=interleukin 2 receptor, gamma chain, precursor /DB_XREF=gi:4557881 /UG=Hs.84 interleukin 2 receptor, gamma (severe combined immunodeficiency) /FL=gb:NM_000206.1	NM_000206		NP_000197

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204122_at	0.016934	gb:NM_003332.1 /DEF=Homo sapiens TYRO protein tyrosine kinase binding protein (TYROBP), mRNA. /FEA=mRNA /GEN=TYROBP /PROD=TYRO protein tyrosine kinase binding protein /DB_XREF=gi:4507754 /UG=Hs.9963 TYRO protein tyrosine kinase binding protein /FL=gb:AF019562.1 gb:NM_003332.1	NM_003332		NP_003323
204125_at	0.008237	gb:NM_016013.1 /DEF=Homo sapiens CGI-65 protein (LOC51103), mRNA. /FEA=mRNA /GEN=LOC51103 /PROD=CGI-65 protein /DB_XREF=gi:7705778 /UG=Hs.106529 CGI-65 protein /FL=gb:BC000780.1 gb:AF151823.1 gb:NM_016013.1	NM_016013		NP_057097
204137_at	0.003368	gb:NM_003272.1 /DEF=Homo sapiens transmembrane 7 superfamily member 1 (upregulated in kidney) (TM7SF1), mRNA. /FEA=mRNA /GEN=TM7SF1 /PROD=transmembrane 7 superfamily member 1(upregulated in kidney) /DB_XREF=gi:4507544 /UG=Hs.15791 transmembrane 7 superfamily member 1 (upregulated in kidney) /FL=gb:AF027826.1 gb:NM_003272.1	NM_003272		NP_003263
204141_at	0.034317	gb:NM_001069.1 /DEF=Homo sapiens tubulin, beta polypeptide (TUBB), mRNA. /FEA=mRNA /GEN=TUBB /PROD=tubulin, beta polypeptide /DB_XREF=gi:4507728 /UG=Hs.179661 tubulin, beta polypeptide /FL=gb:BC001194.1 gb:NM_001069.1	NM_001069		NP_001060

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204150_at	0.013274	gb:NM_015136.1 /DEF=Homo sapiens KIAA0246 protein (KIAA0246), mRNA. /FEA=mRNA /GEN=KIAA0246 /PROD=KIAA0246 protein /DB_XREF=gi:12225239 /UG=Hs.301989 KIAA0246 protein /FL=gb:NM_015136.1	NM_015136		NP_055951
204157_s_at	2.32E-04	gb:NM_025164.1 /DEF=Homo sapiens KIAA0999 protein (KIAA0999), mRNA. /FEA=mRNA /GEN=KIAA0999 /PROD=hypothetical protein FLJ12240 /DB_XREF=gi:13386467 /UG=Hs.4278 KIAA0999 protein /FL=gb:NM_025164.1	NM_025164		NP_079440
204159_at	0.002537	gb:NM_001262.1 /DEF=Homo sapiens cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4) (CDKN2C), mRNA. /FEA=mRNA /GEN=CDKN2C /PROD=cyclin-dependent kinase inhibitor 2C (p18,inhibits CDK4) /DB_XREF=gi:4502750 /UG=Hs.4854 cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4) /FL=gb:AF041248.1 gb:NM_001262.1	NM_001262		NP_523240
204160_s_at	5.03E-04	ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function)	AW194947	Hs.54037	NP_055751
204161_s_at	0.036254	gb:NM_014936.1 /DEF=Homo sapiens ectonucleotide pyrophosphatasephosphodiesterase 4 (putative function) (ENPP4), mRNA. /FEA=mRNA /GEN=ENPP4 /PROD=ectonucleotide pyrophosphatasephosphodiesterase 4 (putative function) /DB_XREF=gi:7662357 /UG=Hs.54037 ectonucleotide pyrophosphatasephosphodiesterase 4 (putative function) /FL=gb:AB020686.1 gb:NM_014936.1	NM_014936		NP_055751

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204168_at	0.035171	gb:NM_002413.1 /DEF=Homo sapiens microsomal glutathione S-transferase 2 (MGST2), mRNA. /FEA=mRNA /GEN=MGST2 /PROD=microsomal glutathione S-transferase 2 /DB_XREF=gi:4505180 /UG=Hs.81874 microsomal glutathione S-transferase 2 /FL=gb:U77604.1 gb:NM_002413.1	NM_002413		NP_002404
204169_at	0.033381	gb:NM_000883.1 /DEF=Homo sapiens IMP (inosine monophosphate) dehydrogenase 1 (IMPDH1), mRNA. /FEA=mRNA /GEN=IMPDH1 /PROD=IMP (inosine monophosphate) dehydrogenase 1 /DB_XREF=gi:4504686 /UG=Hs.850 IMP (inosine monophosphate) dehydrogenase 1 /FL=gb:J05272.1 gb:NM_000883.1	NM_000883		NP_000874
204174_at	0.004341	gb:NM_001629.1 /DEF=Homo sapiens arachidonate 5-lipoxygenase-activating protein (ALOX5AP), mRNA. /FEA=mRNA /GEN=ALOX5AP /PROD=arachidonate 5-lipoxygenase-activating protein /DB_XREF=gi:4502058 /UG=Hs.100194 arachidonate 5-lipoxygenase-activating protein /FL=gb:NM_001629.1	NM_001629		NP_001620
204177_s_at	0.00161	gb:NM_014458.2 /DEF=Homo sapiens Kelch motif containing protein (AB026190), mRNA. /FEA=mRNA /GEN=AB026190 /PROD=Kelch motif containing protein /DB_XREF=gi:13677223 /UG=Hs.106290 Kelch motif containing protein /FL=gb:NM_014458.2 gb:AB026190.1	NM_014458		NP_055273

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204180_s_at	0.007779	Consensus includes gb:AI745225 /FEA=EST /DB_XREF=gi:5113513 /DB_XREF=est:wg10d12.x1 /CLONE=IMAGE:2364695 /UG=Hs.127649 KIAA0414 protein /FL=gb:NM_014007.1	NM_014007		NP_054726
204184_s_at	0.040064	gb:NM_005160.2 /DEF=Homo sapiens adrenergic, beta, receptor kinase 2 (ADRBK2), mRNA. /FEA=mRNA /GEN=ADRBK2 /PROD=beta adrenergic receptor kinase 2 /DB_XREF=gi:6138972 /UG=Hs.13944 adrenergic, beta, receptor kinase 2 /FL=gb:NM_005160.2	NM_005160		NP_005151
204185_x_at	1.26E-04	gb:NM_005038.1 /DEF=Homo sapiens peptidylprolyl isomerase D (cyclophilin D) (PPID), mRNA. /FEA=mRNA /GEN=PPID /PROD=peptidylprolyl isomerase D (cyclophilin D) /DB_XREF=gi:4826931 /UG=Hs.143482 peptidylprolyl isomerase D (cyclophilin D) /FL=gb:L11667.1 gb:NM_005038.1	NM_005038		NP_005029
204186_s_at	0.003171	peptidylprolyl isomerase D (cyclophilin D)	AI014573	Hs.143482	NP_005029
204190_at	0.041055	gb:NM_005800.1 /DEF=Homo sapiens highly charged protein (D13S106E), mRNA. /FEA=mRNA /GEN=D13S106E /PROD=highly charged protein /DB_XREF=gi:5031648 /UG=Hs.151236 highly charged protein /FL=gb:NM_005800.1	NM_005800		NP_005791

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204194_at	0.026891	gb:NM_001186.1 /DEF=Homo sapiens BTB and CNC homology 1, basic leucine zipper transcription factor 1 (BACH1), mRNA. /FEA=mRNA /GEN=BACH1 /PROD=BTB and CNC homology 1, basic leucine zippertranscription factor 1 /DB_XREF=gi:4502352 /UG=Hs.154276 BTB and CNC homology 1, basic leucine zipper transcription factor 1 /FL=gb:AB002803.1 gb:NM_001186.1	NM_001186		NP_001177
204197_s_at	0.041055	gb:NM_004350.1 /DEF=Homo sapiens runt-related transcription factor 3 (RUNX3), mRNA. /FEA=mRNA /GEN=RUNX3 /PROD=runt-related transcription factor 3 /DB_XREF=gi:4757917 /UG=Hs.170019 runt-related transcription factor 3 /FL=gb:NM_004350.1	NM_004350		NP_004341
204198_s_at	5.03E-04	runt-related transcription factor 3	AA541630	Hs.170019	NP_004341
204202_at	0.016934	gb:NM_017604.1 /DEF=Homo sapiens KIAA1023 protein (KIAA1023), mRNA. /FEA=mRNA /GEN=KIAA1023 /PROD=hypothetical protein DKFZp434I0118 /DB_XREF=gi:8922140 /UG=Hs.21361 KIAA1023 protein /FL=gb:NM_017604.1	NM_017604		
204204_at	0.004341	gb:NM_001860.1 /DEF=Homo sapiens solute carrier family 31 (copper transporters), member 2 (SLC31A2), mRNA. /FEA=mRNA /GEN=SLC31A2 /PROD=solute carrier family 31 (copper transporters),member 2 /DB_XREF=gi:4507016 /UG=Hs.24030 solute carrier family 31 (copper transporters), member 2 /FL=gb:U83461.1 gb:NM_001860.1	NM_001860		NP_001851

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204206_at	0.00161	gb:NM_020310.1 /DEF=Homo sapiens MAX binding protein (MNT), mRNA. /FEA=mRNA /GEN=MNT /PROD=MAX binding protein /DB_XREF=gi:9945317 /UG=Hs.25497 MAX binding protein /FL=gb:NM_020310.1	NM_020310		NP_064706
204209_at	0.033381	phosphate cytidyltransferase 1, choline, alpha isoform	AI638771	Hs.273558	
204214_s_at	1.28E-04	gb:NM_006834.1 /DEF=Homo sapiens RAB32, member RAS oncogene family (RAB32), mRNA. /FEA=mRNA /GEN=RAB32 /PROD=RAB32, member RAS oncogene family /DB_XREF=gi:5803132 /UG=Hs.32217 RAB32, member RAS oncogene family /FL=gb:U71127.1 gb:NM_006834.1	NM_006834		NP_006825
204216_s_at	0.012486	gb:NM_024824.1 /DEF=Homo sapiens hypothetical protein FLJ11806 (FLJ11806), mRNA. /FEA=mRNA /GEN=FLJ11806 /PROD=hypothetical protein FLJ11806 /DB_XREF=gi:13376221 /UG=Hs.323443 hypothetical protein FLJ11806 /FL=gb:NM_024824.1	NM_024824		NP_079100
204218_at	1.72E-04	gb:NM_014042.1 /DEF=Homo sapiens DKFZP564M082 protein (DKFZP564M082), mRNA. /FEA=mRNA /GEN=DKFZP564M082 /PROD=DKFZP564M082 protein /DB_XREF=gi:7661621 /UG=Hs.38044 DKFZP564M082 protein /FL=gb:BC005156.1 gb:BC005393.1 gb:AF077206.1 gb:AL080071.1 gb:NM_014042.1	NM_014042		NP_054761

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204219_s_at	3.26E-04	gb:NM_002802.1 /DEF=Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 1 (PSMC1), mRNA. /FEA=mRNA /GEN=PSMC1 /PROD=proteasome (prosome, macropain) 26S subunit,ATPase, 1 /DB_XREF=gi:4506206 /UG=Hs.4745 proteasome (prosome, macropain) 26S subunit, ATPase, 1 /FL=gb:L02426.1 gb:NM_002802.1	NM_002802		NP_002793
204220_at	0.010205	gb:NM_004877.1 /DEF=Homo sapiens glia maturation factor, gamma (GMFG), mRNA. /FEA=mRNA /GEN=GMFG /PROD=glia maturation factor, gamma /DB_XREF=gi:4758439 /UG=Hs.5210 glia maturation factor, gamma /FL=gb:AB001993.1 gb:AF038956.1 gb:NM_004877.1	NM_004877		NP_004868
204225_at	1.26E-04	gb:NM_006037.2 /DEF=Homo sapiens histone deacetylase 4 (HDAC4), mRNA. /FEA=mRNA /GEN=HDAC4 /PROD=histone deacetylase 4 /DB_XREF=gi:13259519 /UG=Hs.91400 histone deacetylase 4 /FL=gb:NM_006037.2 gb:AF132607.1	NM_006037		NP_006028
204226_at	0.00228	gb:NM_014393.1 /DEF=Homo sapiens staufer (Drosophila, RNA binding protein) homolog 2 (STAU2), mRNA. /FEA=mRNA /GEN=STAU2 /PROD=staufer homolog 2 /DB_XREF=gi:7657624 /UG=Hs.96870 staufer (Drosophila, RNA-binding protein) homolog 2 /FL=gb:NM_014393.1	NM_014393		NP_055208

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204227_s_at	0.003171	gb:NM_004614.1 /DEF=Homo sapiens thymidine kinase 2, mitochondrial (TK2), mRNA. /FEA=mRNA /GEN=TK2 /PROD=thymidine kinase 2, mitochondrial /DB_XREF=gi:10281329 /UG=Hs.274701 thymidine kinase 2, mitochondrial /FL=gb:NM_004614.1 gb:U77088.1	NM_004614		NP_004605
204232_at	1.26E-04	gb:NM_004106.1 /DEF=Homo sapiens Fc fragment of IgE, high affinity I, receptor for; gamma polypeptide (FCER1G), mRNA. /FEA=mRNA /GEN=FCER1G /PROD=Fc fragment of IgE, high affinity I, receptorfor, gamma polypeptide precursor /DB_XREF=gi:4758343 /UG=Hs.743 Fc fragment of IgE, high affinity I, receptor for; gamma polypeptide /FL=gb:M33195.1 gb:NM_004106.1	NM_004106		NP_004097
204236_at	1.26E-04	gb:NM_002017.2 /DEF=Homo sapiens Friend leukemia virus integration 1 (FLI1), mRNA. /FEA=mRNA /GEN=FLI1 /PROD=Friend leukemia virus integration 1 /DB_XREF=gi:7110592 /UG=Hs.108043 Friend leukemia virus integration 1 /FL=gb:BC001670.1 gb:M98833.3 gb:NM_002017.2	NM_002017		NP_002008
204246_s_at	0.00161	gb:NM_007234.2 /DEF=Homo sapiens dynactin 3 (p22) (DCTN3), transcript variant 1, mRNA. /FEA=mRNA /GEN=DCTN3 /PROD=dynactin 3, isoform 1 /DB_XREF=gi:13259516 /UG=Hs.15961 dynactin 3 (p22) /FL=gb:NM_007234.2 gb:AF082513.1	NM_007234		NP_077324

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204249_s_at	0.001116	gb:NM_005574.2 /DEF=Homo sapiens LIM domain only 2 (rhombotin-like 1) (LMO2), mRNA. /FEA=mRNA /GEN=LMO2 /PROD=LIM domain only 2 /DB_XREF=gi:6633806 /UG=Hs.184585 LIM domain only 2 (rhombotin-like 1) /FL=gb:NM_005574.2	NM_005574		NP_005565
204252_at	0.026891	gb:M68520.1 /DEF=Human cdc2-related protein kinase mRNA, complete cds. /FEA=mRNA /PROD=cdc2-related protein kinase /DB_XREF=gi:180177 /UG=Hs.19192 cyclin-dependent kinase 2 /FL=gb:BC003065.1 gb:M68520.1 gb:NM_001798.1	M68520		NP_439892
204253_s_at	0.021312	Consensus includes gb:AA454701 /FEA=EST /DB_XREF=gi:2177477 /DB_XREF=est:zx76e01.s1 /CLONE=IMAGE:809688 /UG=Hs.2062 vitamin D (1,25-dihydroxyvitamin D3) receptor /FL=gb:AF026260.1 gb:J03258.1 gb:NM_000376.1	NM_000376		NP_000367
204254_s_at	1.26E-04	gb:NM_000376.1 /DEF=Homo sapiens vitamin D (1,25-dihydroxyvitamin D3) receptor (VDR), mRNA. /FEA=mRNA /GEN=VDR /PROD=vitamin D (1,25- dihydroxyvitamin D3) receptor /DB_XREF=gi:4507882 /UG=Hs.2062 vitamin D (1,25-dihydroxyvitamin D3) receptor /FL=gb:AF026260.1 gb:J03258.1 gb:NM_000376.1	NM_000376		NP_000367
204257_at	0.003862	gb:NM_021727.1 /DEF=Homo sapiens fatty acid desaturase 3 (FADS3), mRNA. /FEA=mRNA /GEN=FADS3 /PROD=fatty acid desaturase 3 /DB_XREF=gi:13375615 /UG=Hs.21765 fatty acid desaturase 3 /FL=gb:AF084560.1 gb:NM_021727.1 gb:BC004901.1 gb:AF134404.1	NM_021727		NP_068373

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204264_at	0.004205	gb:NM_000098.1 /DEF=Homo sapiens carnitine palmitoyltransferase II (CPT2), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=CPT2 /PROD=carnitine palmitoyltransferase II /DB_XREF=gi:4503022 /UG=Hs.274336 carnitine palmitoyltransferase II /FL=gb:U09648.1 gb:BC002445.1 gb:BC005172.1 gb:M58581.1 gb:NM_000098.1	NM_000098		NP_000089
204269_at	0.022795	gb:NM_006875.1 /DEF=Homo sapiens pim-2 oncogene (PIM2), mRNA. /FEA=mRNA /GEN=PIM2 /PROD=pim-2 oncogene /DB_XREF=gi:5803124 /UG=Hs.80205 pim-2 oncogene /FL=gb:U77735.1 gb:NM_006875.1	NM_006875		NP_006866
204275_at	0.001618	small optic lobes homolog (Drosophila)	AI796687	Hs.55836	NP_005623
204287_at	0.034317	gb:NM_004711.1 /DEF=Homo sapiens synaptogyrin 1 (SYNGR1), mRNA. /FEA=mRNA /GEN=SYNGR1 /PROD=synaptogyrin 1 /DB_XREF=gi:4759199 /UG=Hs.6139 synaptogyrin 1 /FL=gb:NM_004711.1	NM_004711		NP_663791
204295_at	0.006096	gb:NM_003172.1 /DEF=Homo sapiens surfait 1 (SURF1), mRNA. /FEA=mRNA /GEN=SURF1 /PROD=surfait 1 /DB_XREF=gi:4507318 /UG=Hs.3196 surfait 1 /FL=gb:NM_003172.1	NM_003172		NP_003163
204297_at	3.26E-04	gb:NM_002647.1 /DEF=Homo sapiens phosphoinositide-3-kinase, class 3 (PIK3C3), mRNA. /FEA=mRNA /GEN=PIK3C3 /PROD=phosphoinositide-3-kinase, class 3 /DB_XREF=gi:4505800 /UG=Hs.32971 phosphoinositide-3-kinase, class 3 /FL=gb:NM_002647.1	NM_002647		NP_002638

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204307_at	0.003862	Consensus includes gb:AB002295.1 /DEF=Human mRNA for KIAA0297 gene, partial cds. /FEA=mRNA /GEN=KIAA0297 /DB_XREF=gi:2224534 /UG=Hs.11711 KIAA0329 gene product /FL=gb:AB002327.1 gb:NM_014844.1	AB002295		NP_055659
204308_s_at	7.58E-04	gb:NM_014844.1 /DEF=Homo sapiens KIAA0329 gene product (KIAA0329), mRNA. /FEA=mRNA /GEN=KIAA0329 /PROD=KIAA0329 gene product /DB_XREF=gi:7662057 /UG=Hs.11711 KIAA0329 gene product /FL=gb:AB002327.1 gb:NM_014844.1	NM_014844		NP_055659
204313_s_at	0.007779	Consensus includes gb:AA161486 /FEA=EST /DB_XREF=gi:1735796 /DB_XREF=est:zq42d09.s1 /CLONE=IMAGE:632369 /UG=Hs.79194 cAMP responsive element binding protein 1 /FL=gb:M27691.1 gb:NM_004379.1	NM_004379		NP_604391
204314_s_at	7.58E-04	gb:NM_004379.1 /DEF=Homo sapiens cAMP responsive element binding protein 1 (CREB1), mRNA. /FEA=mRNA /GEN=CREB1 /PROD=cAMP responsive element binding protein 1 /DB_XREF=gi:4758053 /UG=Hs.79194 cAMP responsive element binding protein 1 /FL=gb:M27691.1 gb:NM_004379.1	NM_004379		NP_604391
204316_at	0.011377	Consensus includes gb:W19676 /FEA=EST /DB_XREF=gi:1295644 /DB_XREF=est:zb36h07.r1 /CLONE=IMAGE:305725 /UG=Hs.82280 regulator of G-protein signalling 10 /FL=gb:NM_002925.2 gb:AF045229.1	NM_002925		NP_002916

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204333_s_at	0.041055	gb:NM_000027.1 /DEF=Homo sapiens aspartylglucosaminidase (AGA), mRNA. /FEA=mRNA /GEN=AGA /PROD=aspartylglucosaminidase precursor /DB_XREF=gi:4557272 /UG=Hs.207776 aspartylglucosaminidase /FL=gb:M64073.1 gb:NM_000027.1	NM_000027		NP_000018
204336_s_at	0.005852	gb:NM_005873.1 /DEF=Homo sapiens regulator of G-protein signalling 19 (RGS19), mRNA. /FEA=mRNA /GEN=RGS19 /PROD=G protein signalling regulator 19 /DB_XREF=gi:5031704 /UG=Hs.22698 regulator of G-protein signalling 19 /FL=gb:NM_005873.1	NM_005873		NP_005864
204342_at	3.26E-04	gb:NM_013386.1 /DEF=Homo sapiens hypothetical protein (DKFZp586G0123), mRNA. /FEA=mRNA /GEN=DKFZp586G0123 /PROD=hypothetical protein /DB_XREF=gi:9558726 /UG=Hs.24713 hypothetical protein /FL=gb:AL050209.1 gb:NM_013386.1	NM_013386		NP_037518
204351_at	0.005852	gb:NM_005980.1 /DEF=Homo sapiens S100 calcium-binding protein P (S100P), mRNA. /FEA=mRNA /GEN=S100P /PROD=S100 calcium-binding protein P /DB_XREF=gi:5174662 /UG=Hs.2962 S100 calcium-binding protein P /FL=gb:NM_005980.1	NM_005980		NP_005971
204361_s_at	0.00228	gb:AB014486.1 /DEF=Homo sapiens mRNA for RA70, complete cds. /FEA=mRNA /GEN=RA70 /PROD=RA70 /DB_XREF=gi:4062959 /UG=Hs.52644 SKAP55 homologue /FL=gb:BC002893.1 gb:AF072166.1 gb:AB014486.1 gb:AF051323.1 gb:NM_003930.1	AB014486		NP_003921

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204362_at	2.05E-04	gb:NM_003930.1 /DEF=Homo sapiens SKAP55 homologue (SKAP-HOM), mRNA. /FEA=mRNA /GEN=SKAP-HOM /PROD=SKAP55 homologue /DB_XREF=gi:4506962 /UG=Hs.52644 SKAP55 homologue /FL=gb:BC002893.1 gb:AF072166.1 gb:AB014486.1 gb:AF051323.1 gb:NM_003930.1	NM_003930		NP_003921
204366_s_at	8.68E-04	gb:NM_001521.1 /DEF=Homo sapiens general transcription factor IIIC, polypeptide 2 (beta subunit, 110kD) (GTF3C2), mRNA. /FEA=mRNA /GEN=GTF3C2 /PROD=general transcription factor IIIC, polypeptide 2(beta subunit, 110kD) /DB_XREF=gi:4504204 /UG=Hs.75782 general transcription factor IIIC, polypeptide 2 (beta subunit, 110kD) /FL=gb:D13636.1 gb:NM_001521.1	NM_001521		NP_001512
204370_at	1.26E-04	gb:NM_006831.1 /DEF=Homo sapiens ATPGTP-binding protein (HEAB), mRNA. /FEA=mRNA /GEN=HEAB /PROD=ATPGTP-binding protein /DB_XREF=gi:5803028 /UG=Hs.87465 ATPGTP-binding protein /FL=gb:BC000446.1 gb:U73524.1 gb:NM_006831.1	NM_006831		NP_006822
204372_s_at	0.033381	gb:NM_003685.1 /DEF=Homo sapiens KH-type splicing regulatory protein (FUSE binding protein 2) (KHSRP), mRNA. /FEA=mRNA /GEN=KHSRP /PROD=KH-type splicing regulatory protein (FUSE binding protein 2) /DB_XREF=gi:4504864 /UG=Hs.91142 KH-type splicing regulatory protein (FUSE binding protein 2) /FL=gb:U94832.1 gb:NM_003685.1	NM_003685		NP_003676

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204373_s_at	0.041055	gb:NM_014810.1 /DEF=Homo sapiens KIAA0480 gene product (KIAA0480), mRNA. /FEA=mRNA /GEN=KIAA0480 /PROD=KIAA0480 gene product /DB_XREF=gi:7662155 /UG=Hs.92200 KIAA0480 gene product /FL=gb:AB007949.1 gb:NM_014810.1	NM_014810		NP_055625
204383_at	0.041605	gb:NM_022719.1 /DEF=Homo sapiens DiGeorge syndrome critical region gene DGS1 (DGS1), mRNA. /FEA=mRNA /GEN=DGS1 /PROD=DiGeorge syndrome critical region gene DGS1protein /DB_XREF=gi:13027629 /UG=Hs.154879 DiGeorge syndrome critical region gene DGS1 /FL=gb:NM_022719.1	NM_022719		NP_073210
204387_x_at	0.005852	gb:NM_024026.1 /DEF=Homo sapiens hypothetical protein MGC3243 (MGC3243), mRNA. /FEA=mRNA /GEN=MGC3243 /PROD=hypothetical protein MGC3243 /DB_XREF=gi:13128969 /UG=Hs.182695 hypothetical protein MGC3243 /FL=gb:BC000002.1 gb:NM_024026.1	NM_024026		NP_076931
204391_x_at	0.004341	gb:NM_015905.1 /DEF=Homo sapiens transcriptional intermediary factor 1 (TIF1), mRNA. /FEA=mRNA /GEN=TIF1 /PROD=transcriptional intermediary factor 1 alpha /DB_XREF=gi:7706233 /UG=Hs.183858 transcriptional intermediary factor 1 /FL=gb:AF009353.1 gb:AF119042.1 gb:NM_003852.1 gb:NM_015905.1	NM_015905		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204393_s_at	2.23E-04	gb:NM_001099.2 /DEF=Homo sapiens acid phosphatase, prostate (ACPP), mRNA. /FEA=mRNA /GEN=ACPP /PROD=prostatic acid phosphatase precursor /DB_XREF=gi:6382063 /UG=Hs.1852 acid phosphatase, prostate /FL=gb:M24902.1 gb:M34840.1 gb:NM_001099.2	NM_001099		NP_001090
204394_at	0.035611	gb:NM_003627.1 /DEF=Homo sapiens prostate cancer overexpressed gene 1 (POV1), mRNA. /FEA=mRNA /GEN=POV1 /PROD=prostate cancer overexpressed gene 1 /DB_XREF=gi:4505970 /UG=Hs.18910 prostate cancer overexpressed gene 1 /FL=gb:BC001639.1 gb:AF045584.1 gb:NM_003627.1	NM_003627		NP_003618
204403_x_at	0.018693	gb:NM_014719.1 /DEF=Homo sapiens KIAA0738 gene product (KIAA0738), mRNA. /FEA=mRNA /GEN=KIAA0738 /PROD=KIAA0738 gene product /DB_XREF=gi:7662275 /UG=Hs.107479 KIAA0738 gene product /FL=gb:AB018281.1 gb:NM_014719.1	NM_014719		NP_055534
204409_s_at	0.001483	gb:BC005248.1 /DEF=Homo sapiens, clone MGC:12282, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:12282) /DB_XREF=gi:13528902 /UG=Hs.155103 eukaryotic translation initiation factor 1A, Y chromosome /FL=gb:BC005248.1 gb:AF000987.1 gb:NM_004681.1	BC005248		NP_004672
204411_at	0.003862	gb:NM_017596.1 /DEF=Homo sapiens KIAA0449 protein (KIAA0449), mRNA. /FEA=mRNA /GEN=KIAA0449 /PROD=hypothetical protein DKFZp434J212 /DB_XREF=gi:8922142 /UG=Hs.169182 KIAA0449 protein /FL=gb:NM_017596.1	NM_017596		NP_060066

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204417_at	1.26E-04	gb:NM_000153.1 /DEF=Homo sapiens galactosylceramidase (Krabbe disease) (GALC), mRNA. /FEA=mRNA /GEN=GALC /PROD=galactosylceramidase precursor /DB_XREF=gi:4557612 /UG=Hs.273 galactosylceramidase (Krabbe disease) /FL=gb:L23116.1 gb:NM_000153.1 gb:D25283.1	NM_000153		NP_000144
204419_x_at	1.72E-04	gb:NM_000184.1 /DEF=Homo sapiens hemoglobin, gamma G (HBG2), mRNA. /FEA=mRNA /GEN=HBG2 /PROD=hemoglobin, gamma G /DB_XREF=gi:6715606 /UG=Hs.283108 hemoglobin, gamma G /FL=gb:NM_000184.1	NM_000184		NP_000175
204423_at	0.034403	gb:NM_013255.1 /DEF=Homo sapiens muskelin 1, intracellular mediator containing kelch motifs (MKLN1), mRNA. /FEA=mRNA /GEN=MKLN1 /PROD=muskelin 1, intracellular mediator containing kelch motifs /DB_XREF=gi:7019458 /UG=Hs.288791 muskelin 1, intracellular mediator containing kelch motifs /FL=gb:BC002834.1 gb:AF047489.1 gb:NM_013255.1	NM_013255		NP_037387
204427_s_at	0.033491	gb:NM_006815.1 /DEF=Homo sapiens coated vesicle membrane protein (RNP24), mRNA. /FEA=mRNA /GEN=RNP24 /PROD=coated vesicle membrane protein /DB_XREF=gi:5803148 /UG=Hs.323378 coated vesicle membrane protein /FL=gb:NM_006815.1	NM_006815		NP_006806
204445_s_at	0.005039	arachidonate 5-lipoxygenase	AI361850	Hs.89499	NP_000689

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204446_s_at	0.00161	gb:NM_000698.1 /DEF=Homo sapiens arachidonate 5-lipoxygenase (ALOX5), mRNA. /FEA=mRNA /GEN=ALOX5 /PROD=arachidonate 5-lipoxygenase /DB_XREF=gi:4502056 /UG=Hs.89499 arachidonate 5-lipoxygenase /FL=gb:J03600.1 gb:J03571.1 gb:NM_000698.1	NM_000698		NP_000689
204461_x_at	0.033381	gb:NM_002853.1 /DEF=Homo sapiens RAD1 (S. pombe) homolog (RAD1), mRNA. /FEA=mRNA /GEN=RAD1 /PROD=RAD1 (S. pombe) homolog /DB_XREF=gi:4506384 /UG=Hs.7179 RAD1 (S. pombe) homolog /FL=gb:AF058392.1 gb:AF073524.1 gb:AF074717.1 gb:AF011905.1 gb:AF084512.1 gb:AF030933.1 gb:AF076841.1 gb:NM_002853.1	NM_002853		NP_596868
204466_s_at	1.26E-04	synuclein, alpha (non A4 component of amyloid precursor)	BG260394	Hs.76930	NP_009292
204467_s_at	0.001053	gb:NM_000345.2 /DEF=Homo sapiens synuclein, alpha (non A4 component of amyloid precursor) (SNCA), transcript variant NACP140, mRNA. /FEA=mRNA /GEN=SNCA /PROD=alpha-synuclein isoform NACP140 /DB_XREF=gi:6806896 /UG=Hs.76930 synuclein, alpha (non A4 component of amyloid precursor) /FL=gb:L08850.1 gb:NM_000345.2	NM_000345		NP_009292
204478_s_at	1.28E-04	gb:NM_002871.1 /DEF=Homo sapiens RAB interacting factor (RABIF), mRNA. /FEA=mRNA /GEN=RABIF /PROD=RAB interacting factor /DB_XREF=gi:4506378 /UG=Hs.90875 RAB interacting factor /FL=gb:U74324.1 gb:NM_002871.1	NM_002871		NP_002862

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204489_s_at	0.003171	gb:NM_000610.1 /DEF=Homo sapiens CD44 antigen (homing function and Indian blood group system) (CD44), mRNA. /FEA=mRNA /GEN=CD44 /PROD=CD44 antigen (homing function and Indian bloodgroup system) /DB_XREF=gi:10835162 /UG=Hs.169610 CD44 antigen (homing function and Indian blood group system) /FL=gb:NM_000610.1 gb:U40373.1 gb:M59040.1 gb:M24915.1	NM_000610		NP_000601
204491_at	0.013223	phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 duncce homolog, Drosophila)	R40917	Hs.172081	NP_006194
204496_at	0.00228	gb:NM_014574.1 /DEF=Homo sapiens nuclear autoantigen (GS2NA), mRNA. /FEA=mRNA /GEN=GS2NA /PROD=nuclear autoantigen /DB_XREF=gi:11128016 /UG=Hs.183105 nuclear autoantigen /FL=gb:NM_014574.1 gb:U17989.1	NM_014574		NP_055389
204500_s_at	1.26E-04	gb:NM_015239.1 /DEF=Homo sapiens KIAA1035 protein (KIAA1035), mRNA. /FEA=mRNA /GEN=KIAA1035 /PROD=KIAA1035 protein /DB_XREF=gi:7662451 /UG=Hs.21542 KIAA1035 protein /FL=gb:NM_015239.1	NM_015239		NP_056054
204505_s_at	1.26E-04	gb:NM_001978.1 /DEF=Homo sapiens erythrocyte membrane protein band 4.9 (dematin) (EPB49), mRNA. /FEA=mRNA /GEN=EPB49 /PROD=erythrocyte membrane protein band 4.9 (dematin) /DB_XREF=gi:4503580 /UG=Hs.274122 erythrocyte membrane protein band 4.9 (dematin) /FL=gb:NM_001978.1 gb:U28389.1	NM_001978		NP_001969

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204507_s_at	0.016934	gb:NM_000945.1 /DEF=Homo sapiens protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcineurin B, type I) (PPP3R1), mRNA. /FEA=mRNA /GEN=PPP3R1 /PROD=protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcineurin B, type I) /DB_XREF=gi:4506024 /UG=Hs.278540 protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcineurin B, type I) /FL=gb:M30773.1 gb:NM_000945.1	NM_000945		NP_000936
204513_s_at	1.26E-04	gb:NM_014800.1 /DEF=Homo sapiens KIAA0281 gene product (KIAA0281), mRNA. /FEA=mRNA /GEN=KIAA0281 /PROD=KIAA0281 gene product /DB_XREF=gi:7662037 /UG=Hs.31463 KIAA0281 gene product /FL=gb:BC003051.1 gb:D87457.1 gb:NM_014800.1	NM_014800		NP_569709
204520_x_at	0.033381	gb:NM_014577.1 /DEF=Homo sapiens bromodomain-containing 1 (BRD1), mRNA. /FEA=mRNA /GEN=BRD1 /PROD=bromodomain-containing protein 1 /DB_XREF=gi:11321641 /UG=Hs.127950 bromodomain-containing 1 /FL=gb:NM_014577.1 gb:AF005067.1	NM_014577		NP_055392
204525_at	0.023856	gb:NM_014660.1 /DEF=Homo sapiens KIAA0783 gene product (KIAA0783), mRNA. /FEA=mRNA /GEN=KIAA0783 /PROD=KIAA0783 gene product /DB_XREF=gi:7662303 /UG=Hs.156276 KIAA0783 gene product /FL=gb:AB018326.1 gb:NM_014660.1	NM_014660		NP_055475

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204526_s_at	1.26E-04	gb:NM_007063.1 /DEF=Homo sapiens vascular Rab-GAPTBC-containing (VRP), mRNA. /FEA=mRNA /GEN=VRP /PROD=vascular Rab-GAPTBC-containing /DB_XREF=gi:5902153 /UG=Hs.164170 vascular Rab-GAPTBC-containing /FL=gb:AB024057.1 gb:NM_007063.1	NM_007063		NP_008994
204527_at	0.003171	gb:NM_000259.1 /DEF=Homo sapiens myosin VA (heavy polypeptide 12, myoxin) (MYO5A), mRNA. /FEA=mRNA /GEN=MYO5A /PROD=myosin VA (heavy polypeptide 12, myoxin) /DB_XREF=gi:10835118 /UG=Hs.170157 myosin VA (heavy polypeptide 12, myoxin) /FL=gb:NM_000259.1 gb:U90942.1	NM_000259		NP_000250
204531_s_at	0.03878	gb:NM_007295.1 /DEF=Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1b, mRNA. /FEA=mRNA /GEN=BRCA1 /PROD=breast cancer 1, early onset /DB_XREF=gi:6552300 /UG=Hs.194143 breast cancer 1, early onset /FL=gb:NM_007295.1	NM_007295		NP_009237
204542_at	0.042536	gb:NM_006456.1 /DEF=Homo sapiens sialyltransferase (STHM), mRNA. /FEA=mRNA /GEN=STHM /PROD=sialyltransferase /DB_XREF=gi:5454091 /UG=Hs.288215 sialyltransferase /FL=gb:NM_006456.1 gb:U14550.1	NM_006456		NP_006447
204546_at	1.26E-04	gb:NM_014732.1 /DEF=Homo sapiens KIAA0513 gene product (KIAA0513), mRNA. /FEA=mRNA /GEN=KIAA0513 /PROD=KIAA0513 gene product /DB_XREF=gi:7662163 /UG=Hs.301658 KIAA0513 gene product /FL=gb:AB011085.1 gb:NM_014732.1	NM_014732		NP_055547

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204554_at	1.28E-04	Consensus includes gb:AL109928 /DEF=Human DNA sequence from clone RP4-551D2 on chromosome 20q13.2-13.33. Contains the gene for a novel Cadherin domain protein, a novel gene, the PPP1R6 gene for protein phosphatase 1 regulatory subunit 6, the 5 end of the SYCP2 gene for synaptonemal... /FEA=mRNA_6 /DB_XREF=gi:7981303 /UG=Hs.42215 protein phosphatase 1, regulatory subunit 6 /FL=gb:NM_006242.2	AL109928		
204565_at	1.28E-04	gb:NM_018473.1 /DEF=Homo sapiens uncharacterized hypothalamus protein HT012 (HT012), mRNA. /FEA=mRNA /GEN=HT012 /PROD=uncharacterized hypothalamus protein HT012 /DB_XREF=gi:8923811 /UG=Hs.9676 uncharacterized hypothalamus protein HT012 /FL=gb:BC000894.1 gb:AF274952.1 gb:AF220186.1 gb:NM_018473.1	NM_018473		NP_060943
204593_s_at	0.033381	hypothetical protein FLJ20232	AA046752	Hs.83869	NP_037430
204599_s_at	0.001551	gb:NM_006428.1 /DEF=Homo sapiens melanoma-associated antigen recognised by cytotoxic T lymphocytes (MAAT1), mRNA. /FEA=mRNA /GEN=MAAT1 /PROD=melanoma-associated antigen recognised by Tlymphocytes /DB_XREF=gi:5453835 /UG=Hs.279869 melanoma-associated antigen recognised by cytotoxic T lymphocytes /FL=gb:BC000507.1 gb:BC000990.2 gb:NM_006428.1 gb:U19796.1	NM_006428		NP_006419

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204604_at	2.05E-04	gb:NM_012395.1 /DEF=Homo sapiens PFTAIR protein kinase 1 (PFTK1), mRNA. /FEA=mRNA /GEN=PFTK1 /PROD=PFTAIR protein kinase 1 /DB_XREF=gi:6912583 /UG=Hs.57856 PFTAIR protein kinase 1 /FL=gb:AB020641.1 gb:NM_012395.1	NM_012395		NP_036527
204613_at	5.03E-04	gb:NM_002661.1 /DEF=Homo sapiens phospholipase C, gamma 2 (phosphatidylinositol-specific) (PLCG2), mRNA. /FEA=mRNA /GEN=PLCG2 /PROD=phospholipase C, gamma 2(phosphatidylinositol-specific) /DB_XREF=gi:4505870 /UG=Hs.75648 phospholipase C, gamma 2 (phosphatidylinositol-specific) /FL=gb:M37238.1 gb:NM_002661.1	NM_002661		NP_002652
204614_at	1.58E-04	gb:NM_002575.1 /DEF=Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2 (SERPINB2), mRNA. /FEA=mRNA /GEN=SERPINB2 /PROD=serine (or cysteine) proteinase inhibitor, cladeB (ovalbumin), member 2 /DB_XREF=gi:4505594 /UG=Hs.75716 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2 /FL=gb:J02685.1 gb:J03603.1 gb:M18082.1 gb:NM_002575.1	NM_002575		NP_002566
204615_x_at	1.26E-04	gb:NM_004508.1 /DEF=Homo sapiens isopentenyl-diphosphate delta isomerase (IDI1), mRNA. /FEA=mRNA /GEN=IDI1 /PROD=isopentenyl-diphosphate delta isomerase /DB_XREF=gi:4758583 /UG=Hs.76038 isopentenyl-diphosphate delta isomerase /FL=gb:NM_004508.1	NM_004508		NP_004499

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204616_at	0.004644	gb:NM_006002.1 /DEF=Homo sapiens ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase) (UCHL3), mRNA. /FEA=mRNA /GEN=UCHL3 /PROD=ubiquitin carboxyl-terminal esterase L3(ubiquitin thiolesterase) /DB_XREF=gi:5174740 /UG=Hs.77917 ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase) /FL=gb:M30496.1 gb:NM_006002.1	NM_006002		NP_005993
204619_s_at	0.00161	chondroitin sulfate proteoglycan 2 (versican)	BF590263	Hs.81800	NP_004376
204620_s_at	1.26E-04	gb:NM_004385.1 /DEF=Homo sapiens chondroitin sulfate proteoglycan 2 (versican) (CSPG2), mRNA. /FEA=mRNA /GEN=CSPG2 /PROD=chondroitin sulfate proteoglycan 2 (versican) /DB_XREF=gi:4758081 /UG=Hs.81800 chondroitin sulfate proteoglycan 2 (versican) /FL=gb:NM_004385.1	NM_004385		NP_004376
204630_s_at	0.005039	gb:NM_004871.1 /DEF=Homo sapiens golgi SNAP receptor complex member 1 (GOSR1), mRNA. /FEA=mRNA /GEN=GOSR1 /PROD=golgi SNAP receptor complex member 1 /DB_XREF=gi:4758455 /UG=Hs.8868 golgi SNAP receptor complex member 1 /FL=gb:AF073926.1 gb:NM_004871.1	NM_004871		NP_004862

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204632_at	0.003171	gb:NM_003942.1 /DEF=Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 4 (RPS6KA4), mRNA. /FEA=mRNA /GEN=RPS6KA4 /PROD=ribosomal protein S6 kinase, 90kD, polypeptide4 /DB_XREF=gi:4506734 /UG=Hs.105584 ribosomal protein S6 kinase, 90kD, polypeptide 4 /FL=gb:NM_003942.1	NM_003942		NP_003933
204633_s_at	0.005039	gb:AF074393.1 /DEF=Homo sapiens nuclear mitogen- and stress-activated protein kinase-1 (MSK1) mRNA, complete cds. /FEA=mRNA /GEN=MSK1 /PROD=nuclear mitogen- and stress-activated proteinkinase-1 /DB_XREF=gi:3411156 /UG=Hs.109058 ribosomal protein S6 kinase, 90kD, polypeptide 5 /FL=gb:AF074393.1 gb:AF090421.1 gb:AF080000.1 gb:NM_004755.1	AF074393		NP_004746
204640_s_at	0.026891	gb:NM_003563.1 /DEF=Homo sapiens speckle-type POZ protein (SPOP), mRNA. /FEA=mRNA /GEN=SPOP /PROD=speckle-type POZ protein /DB_XREF=gi:4507182 /UG=Hs.129951 speckle-type POZ protein /FL=gb:BC003385.1 gb:NM_003563.1	NM_003563		NP_003554
204646_at	7.58E-04	gb:NM_000110.2 /DEF=Homo sapiens dihydropyrimidine dehydrogenase (DPYD), mRNA. /FEA=mRNA /GEN=DPYD /PROD=dihydropyrimidine dehydrogenase /DB_XREF=gi:4557874 /UG=Hs.1602 dihydropyrimidine dehydrogenase /FL=gb:U20938.1 gb:NM_000110.2 gb:U09178.1 gb:AB003063.1	NM_000110		NP_000101

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204647_at	0.033381	gb:NM_004838.1 /DEF=Homo sapiens Homer, neuronal immediate early gene, 3 (HOMER-3), mRNA. /FEA=mRNA /GEN=HOMER-3 /PROD=Homer, neuronal immediate early gene, 3 /DB_XREF=gi:4758549 /UG=Hs.166146 Homer, neuronal immediate early gene, 3 /FL=gb:AF093265.1 gb:NM_004838.1	NM_004838		NP_004829
204651_at	0.003171	nuclear respiratory factor 1	AW003022	Hs.180069	
204652_s_at	0.016934	gb:NM_005011.1 /DEF=Homo sapiens nuclear respiratory factor 1 (NRF1), mRNA. /FEA=mRNA /GEN=NRF1 /PROD=nuclear respiratory factor 1 /DB_XREF=gi:4826865 /UG=Hs.180069 nuclear respiratory factor 1 /FL=gb:U02683.1 gb:NM_005011.1	NM_005011		NP_005002
204653_at	0.041795	transcription factor AP-2 alpha (activating enhancer binding protein 2 alpha)	BF343007	Hs.334334	NP_003211
204661_at	1.26E-04	gb:NM_001803.1 /DEF=Homo sapiens CDW52 antigen (CAMPATH-1 antigen) (CDW52), mRNA. /FEA=mRNA /GEN=CDW52 /PROD=CDW52 antigen (CAMPATH-1 antigen) /DB_XREF=gi:4502760 /UG=Hs.276770 CDW52 antigen (CAMPATH-1 antigen) /FL=gb:BC000644.1 gb:NM_001803.1	NM_001803		NP_001794
204669_s_at	0.007258	gb:NM_007219.2 /DEF=Homo sapiens ring finger protein 24 (RNF24), mRNA. /FEA=mRNA /GEN=RNF24 /PROD=ring finger protein 24 /DB_XREF=gi:10518498 /UG=Hs.30524 ring finger protein 24 /FL=gb:NM_007219.2 gb:AF151081.1	NM_007219		NP_009150

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204672_s_at	0.045541	gb:NM_014942.1 /DEF=Homo sapiens KIAA0957 protein (KIAA0957), mRNA. /FEA=mRNA /GEN=KIAA0957 /PROD=KIAA0957 protein /DB_XREF=gi:7662405 /UG=Hs.30991 KIAA0957 protein /FL=gb:AB023174.1 gb:NM_014942.1	NM_014942		NP_055757
204682_at	0.00424	gb:NM_000428.1 /DEF=Homo sapiens latent transforming growth factor beta binding protein 2 (LTBP2), mRNA. /FEA=mRNA /GEN=LTBP2 /PROD=latent transforming growth factor beta bindingprotein 2 /DB_XREF=gi:4557732 /UG=Hs.83337 latent transforming growth factor beta binding protein 2 /FL=gb:NM_000428.1	NM_000428		NP_000419
204683_at	0.001116	gb:NM_000873.2 /DEF=Homo sapiens intercellular adhesion molecule 2 (ICAM2), mRNA. /FEA=mRNA /GEN=ICAM2 /PROD=intercellular adhesion molecule 2 precursor /DB_XREF=gi:12545398 /UG=Hs.83733 intercellular adhesion molecule 2 /FL=gb:NM_000873.2 gb:BC003097.1	NM_000873		NP_000864
204689_at	5.03E-04	gb:NM_001529.1 /DEF=Homo sapiens hematopoietically expressed homeobox (HHEX), mRNA. /FEA=mRNA /GEN=HHEX /PROD=hematopoietically expressed homeobox /DB_XREF=gi:10835016 /UG=Hs.118651 hematopoietically expressed homeobox /FL=gb:NM_001529.1 gb:L16499.1 gb:NM_002729.1	NM_001529		

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
204690_at	0.007779	gb:NM_004853.1 /DEF=Homo sapiens syntaxin 8 (STX8), mRNA. /FEA=mRNA /GEN=STX8 /PROD=syntaxin 8 /DB_XREF=gi:4759187 /UG=Hs.119525 syntaxin 8 /FL=gb:AF062077.1 gb:AF036715.1 gb:AF115323.1 gb:NM_004853.1	NM_004853		NP_004844
204714_s_at	1.26E-04	gb:NM_000130.2 /DEF=Homo sapiens coagulation factor V (proaccelerin, labile factor) (F5), mRNA. /FEA=mRNA /GEN=F5 /PROD=coagulation factor V precursor /DB_XREF=gi:10518500 /UG=Hs.30054 coagulation factor V (proaccelerin, labile factor) /FL=gb:NM_000130.2 gb:M16967.1 gb:M14335.1	NM_000130		NP_000121
204725_s_at	0.042536	gb:NM_006153.1 /DEF=Homo sapiens NCK adaptor protein 1 (NCK1), mRNA. /FEA=mRNA /GEN=NCK1 /PROD=NCK adaptor protein 1 /DB_XREF=gi:5453753 /UG=Hs.54589 NCK adaptor protein 1 /FL=gb:NM_006153.1	NM_006153		NP_006144
204732_s_at	0.003529	ADP-ribosylation factor domain protein 1, 64kDa	AI021991	Hs.792	NP_150231
204759_at	0.016934	gb:NM_001268.1 /DEF=Homo sapiens chromosome condensation 1-like (CHC1L), mRNA. /FEA=mRNA /GEN=CHC1L /PROD=RCC1-like G exchanging factor RLG /DB_XREF=gi:4557444 /UG=Hs.27007 chromosome condensation 1-like /FL=gb:AF060219.1 gb:NM_001268.1	NM_001268		NP_001259

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204767_s_at	0.001198	gb:BC000323.1 /DEF=Homo sapiens, flap structure-specific endonuclease 1, clone MGC:8478, mRNA, complete cds. /FEA=mRNA /PROD=flap structure-specific endonuclease 1 /DB_XREF=gi:12653112 /UG=Hs.4756 flap structure-specific endonuclease 1 /FL=gb:BC000323.1 gb:NM_004111.3	BC000323		NP_004102
204771_s_at	6.68E-04	transcription termination factor, RNA polymerase I	AI632304	Hs.54780	NP_031370
204772_s_at	0.013088	gb:NM_007344.1 /DEF=Homo sapiens transcription termination factor, RNA polymerase I (TTF1), mRNA. /FEA=mRNA /GEN=TTF1 /PROD=transcription termination factor, RNA polymerase I /DB_XREF=gi:6678454 /UG=Hs.54780 transcription termination factor, RNA polymerase I /FL=gb:NM_007344.1	NM_007344		NP_031370
204786_s_at	0.005411	gb:L41944.1 /DEF=Homo sapiens interferon receptor ifnar2-1 (splice variant IFNAR2-1) mRNA, complete cds. /FEA=mRNA /GEN=IFNAR2 /PROD=interferon receptor /DB_XREF=gi:995296 /UG=Hs.86958 interferon (alpha, beta and omega) receptor 2 /FL=gb:NM_000874.1 gb:L41944.1	L41944		NP_000865
204798_at	8.94E-04	gb:NM_005375.1 /DEF=Homo sapiens v-myb avian myeloblastosis viral oncogene homolog (MYB), mRNA. /FEA=mRNA /GEN=MYB /PROD=v-myb avian myeloblastosis viral oncogene homolog /DB_XREF=gi:4885496 /UG=Hs.1334 v-myb avian myeloblastosis viral oncogene homolog /FL=gb:M15024.1 gb:AF104863.1 gb:NM_005375.1	NM_005375		NP_005366
204800_s_at	0.005852	hypothetical protein FLJ13639	AW194384	Hs.101821	NP_078981

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204806_x_at	0.02145	gb:NM_018950.1 /DEF=Homo sapiens major histocompatibility complex, class I, F (HLA-F), mRNA. /FEA=mRNA /GEN=HLA-F /PROD=major histocompatibility complex, class I, F /DB_XREF=gi:9665231 /UG=Hs.110309 major histocompatibility complex, class I, F /FL=gb:NM_018950.1	NM_018950		NP_061823
204808_s_at	0.020412	gb:NM_014254.1 /DEF=Homo sapiens transmembrane protein 5 (TMEM5), mRNA. /FEA=mRNA /GEN=TMEM5 /PROD=transmembrane protein 5 /DB_XREF=gi:7657177 /UG=Hs.112986 transmembrane protein 5 /FL=gb:AB015633.1 gb:NM_014254.1	NM_014254		NP_055069
204809_at	0.010205	gb:NM_006660.3 /DEF=Homo sapiens ClpX (caseinolytic protease X, E. coli) homolog (CLPX), mRNA. /FEA=mRNA /GEN=CLPX /PROD=ClpX (caseinolytic protease X, E. coli) homolog /DB_XREF=gi:12597621 /UG=Hs.113823 ClpX (caseinolytic protease X, E. coli) homolog /FL=gb:NM_006660.3	NM_006660		NP_006651
204820_s_at	0.016934	gb:NM_006994.2 /DEF=Homo sapiens butyrophilin, subfamily 3, member A3 (BTN3A3), mRNA. /FEA=mRNA /GEN=BTN3A3 /PROD=butyrophilin, subfamily 3, member A3 /DB_XREF=gi:6325463 /UG=Hs.167741 butyrophilin, subfamily 3, member A3 /FL=gb:U90548.1 gb:NM_006994.2	NM_006994		NP_008925

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204821_at	0.026891	gb:NM_006994.2 /DEF=Homo sapiens butyrophilin, subfamily 3, member A3 (BTN3A3), mRNA. /FEA=mRNA /GEN=BTN3A3 /PROD=butyrophilin, subfamily 3, member A3 /DB_XREF=gi:6325463 /UG=Hs.167741 butyrophilin, subfamily 3, member A3 /FL=gb:U90548.1 gb:NM_006994.2	NM_006994		NP_008925
204837_at	0.004341	Consensus includes gb:AL080178.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434K171 (from clone DKFZp434K171); partial cds. /FEA=mRNA /GEN=DKFZp434K171 /PROD=hypothetical protein /DB_XREF=gi:5262652 /UG=Hs.27194 DKFZP434K171 protein /FL=gb:NM_015458.1	AL080178		NP_056273
204848_x_at	0.034317	gb:NM_000559.1 /DEF=Homo sapiens hemoglobin, gamma A (HBG1), mRNA. /FEA=mRNA /GEN=HBG1 /PROD=hemoglobin, gamma A /DB_XREF=gi:4504352 /UG=Hs.305960 hemoglobin, gamma A /FL=gb:AF130098.1 gb:NM_000559.1	NM_000559		NP_000550
204849_at	5.03E-04	gb:NM_006602.1 /DEF=Homo sapiens transcription factor-like 5 (basic helix-loop-helix) (TCFL5), mRNA. /FEA=mRNA /GEN=TCFL5 /PROD=transcription factor-like 5 (basic helix-loop-helix) /DB_XREF=gi:5730082 /UG=Hs.30696 transcription factor-like 5 (basic helix-loop-helix) /FL=gb:AB012124.1 gb:NM_006602.1 gb:AF070992.1	NM_006602		NP_006593

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204859_s_at	7.58E-04	gb:NM_013229.1 /DEF=Homo sapiens apoptotic protease activating factor (APAF1), transcript variant 1, mRNA. /FEA=mRNA /GEN=APAF1 /PROD=apoptotic protease activating factor isoform a /DB_XREF=gi:7108332 /UG=Hs.77579 apoptotic protease activating factor /FL=gb:AB007873.1 gb:AF134397.1 gb:NM_013229.1	NM_013229		NP_037361
204860_s_at	0.016934	baculoviral IAP repeat-containing 1	AI817801	Hs.79019	NP_004527
204861_s_at	1.26E-04	gb:NM_004536.1 /DEF=Homo sapiens baculoviral IAP repeat-containing 1 (BIRC1), mRNA. /FEA=mRNA /GEN=BIRC1 /PROD=baculoviral IAP repeat-containing 1 /DB_XREF=gi:4758751 /UG=Hs.79019 baculoviral IAP repeat-containing 1 /FL=gb:U19251.1 gb:NM_004536.1	NM_004536		NP_004527
204875_s_at	0.031948	gb:NM_001500.1 /DEF=Homo sapiens GDP-mannose 4,6-dehydratase (GMDS), mRNA. /FEA=mRNA /GEN=GMDS /PROD=GDP-mannose 4,6-dehydratase /DB_XREF=gi:4504030 /UG=Hs.105435 GDP-mannose 4,6-dehydratase /FL=gb:BC000117.1 gb:AF042377.1 gb:NM_001500.1	NM_001500		NP_001491
204881_s_at	0.04734	gb:NM_003358.1 /DEF=Homo sapiens UDP-glucose ceramide glucosyltransferase (UGCG), mRNA. /FEA=mRNA /GEN=UGCG /PROD=ceramide glucosyltransferase /DB_XREF=gi:4507810 /UG=Hs.152601 UDP-glucose ceramide glucosyltransferase /FL=gb:D50840.1 gb:NM_003358.1	NM_003358		NP_003349

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204890_s_at	6.68E-04	gb:U07236.1 /DEF=Human mutant lymphocyte-specific protein tyrosine kinase (LCK) mRNA, complete cds. /FEA=mRNA /GEN=LCK /PROD=lymphocyte-specific protein tyrosine kinase /DB_XREF=gi:460965 /UG=Hs.1765 lymphocyte-specific protein tyrosine kinase /FL=gb:M36881.1 gb:U07236.1 gb:NM_005356.1	U07236		NP_005347
204891_s_at	0.004341	gb:NM_005356.1 /DEF=Homo sapiens lymphocyte-specific protein tyrosine kinase (LCK), mRNA. /FEA=mRNA /GEN=LCK /PROD=lymphocyte-specific protein tyrosine kinase /DB_XREF=gi:4885448 /UG=Hs.1765 lymphocyte-specific protein tyrosine kinase /FL=gb:M36881.1 gb:U07236.1 gb:NM_005356.1	NM_005356		NP_005347
204892_x_at	0.02145	gb:NM_001402.1 /DEF=Homo sapiens eukaryotic translation elongation factor 1 alpha 1 (EEF1A1), mRNA. /FEA=mRNA /GEN=EEF1A1 /PROD=eukaryotic translation elongation factor 1 alpha1 /DB_XREF=gi:4503470 /UG=Hs.181165 eukaryotic translation elongation factor 1 alpha 1 /FL=gb:AF267861.1 gb:NM_001402.1	NM_001402		NP_001393
204923_at	0.016934	Consensus includes gb:AL023653 /DEF=Human DNA sequence from clone 753P9 on chromosome Xq25-26.1. Contains the gene coding for Aminopeptidase P (EC 3.4.11.9, XAA-ProX-ProProlineAminoacylproline Aminopeptidase) and a novel gene. Contains ESTs, STSs, GSSs and a gaaa repeat polymorphism /FEA=mRNA_1 /DB_XREF=gi:3550108 /UG=Hs.61469 hypothetical protein /FL=gb:NM_018990.1	AL023653		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
204924_at	0.00228	gb:NM_003264.1 /DEF=Homo sapiens toll-like receptor 2 (TLR2), mRNA. /FEA=mRNA /GEN=TLR2 /PROD=toll-like receptor2 /DB_XREF=gi:4507528 /UG=Hs.63668 toll-like receptor 2 /FL=gb:U88878.1 gb:AF051152.1 gb:NM_003264.1	NM_003264		NP_003255
204971_at	0.004341	gb:NM_005213.1 /DEF=Homo sapiens cystatin A (stefin A) (CSTA), mRNA. /FEA=mRNA /GEN=CSTA /PROD=cystatin A (stefin A) /DB_XREF=gi:4885164 /UG=Hs.2621 cystatin A (stefin A) /FL=gb:NM_005213.1	NM_005213		NP_005204
205003_at	0.001116	gb:NM_014705.1 /DEF=Homo sapiens KIAA0716 gene product (KIAA0716), mRNA. /FEA=mRNA /GEN=KIAA0716 /PROD=KIAA0716 gene product /DB_XREF=gi:7662263 /UG=Hs.118140 KIAA0716 gene product /FL=gb:AB018259.1 gb:NM_014705.1	NM_014705		NP_055520
205010_at	0.027418	gb:NM_019067.1 /DEF=Homo sapiens hypothetical protein (FLJ10613), mRNA. /FEA=mRNA /GEN=FLJ10613 /PROD=hypothetical protein /DB_XREF=gi:9506610 /UG=Hs.142217 hypothetical protein /FL=gb:NM_019067.1	NM_019067		NP_061940
205011_at	0.001078	gb:NM_014622.1 /DEF=Homo sapiens loss of heterozygosity, 11, chromosomal region 2, gene A (LOH11CR2A), mRNA. /FEA=mRNA /GEN=LOH11CR2A /PROD=loss of heterozygosity, 11, chromosomal region2, gene A /DB_XREF=gi:7657310 /UG=Hs.152944 loss of heterozygosity, 11, chromosomal region 2, gene A /FL=gb:AF002672.1 gb:NM_014622.1	NM_014622		NP_055437

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205012_s_at	0.038554	gb:NM_005326.1 /DEF=Homo sapiens hydroxyacyl glutathione hydrolase (HAGH), mRNA. /FEA=mRNA /GEN=HAGH /PROD=hydroxyacyl glutathione hydrolase /DB_XREF=gi:4885388 /UG=Hs.155482 hydroxyacyl glutathione hydrolase /FL=gb:BC000840.1 gb:BC002627.1 gb:NM_005326.1	NM_005326		NP_005317
205013_s_at	7.58E-04	gb:NM_000675.2 /DEF=Homo sapiens adenosine A2a receptor (ADORA2A), mRNA. /FEA=mRNA /GEN=ADORA2A /PROD=adenosine A2a receptor /DB_XREF=gi:5921991 /UG=Hs.1613 adenosine A2a receptor /FL=gb:M97370.1 gb:NM_000675.2	NM_000675		NP_000666
205016_at	0.007779	gb:NM_003236.1 /DEF=Homo sapiens transforming growth factor, alpha (TGFA), mRNA. /FEA=mRNA /GEN=TGFA /PROD=transforming growth factor, alpha /DB_XREF=gi:4507460 /UG=Hs.170009 transforming growth factor, alpha /FL=gb:BC005308.1 gb:M31172.1 gb:K03222.1 gb:NM_003236.1	NM_003236		NP_003227
205021_s_at	0.001205	checkpoint suppressor 1	AA860806	Hs.211773	NP_005188
205027_s_at	5.03E-04	gb:NM_005204.1 /DEF=Homo sapiens mitogen-activated protein kinase kinase 8 (MAP3K8), mRNA. /FEA=mRNA /GEN=MAP3K8 /PROD=mitogen-activated protein kinase kinase 8 /DB_XREF=gi:4885146 /UG=Hs.248 mitogen-activated protein kinase kinase 8 /FL=gb:D14497.1 gb:NM_005204.1	NM_005204		NP_005195

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205033_s_at	7.58E-04	gb:NM_004084.2 /DEF=Homo sapiens defensin, alpha 1, myeloid-related sequence (DEFA1), mRNA. /FEA=mRNA /GEN=DEFA1 /PROD=defensin, alpha 1, preproprotein /DB_XREF=gi:12621915 /UG=Hs.274463 defensin, alpha 1, myeloid-related sequence /FL=gb:NM_004084.2 gb:M21130.1 gb:M26602.1	NM_004084		NP_004075
205034_at	0.001053	gb:NM_004702.1 /DEF=Homo sapiens cyclin E2 (CCNE2), mRNA. /FEA=mRNA /GEN=CCNE2 /PROD=cyclin E2 /DB_XREF=gi:4757931 /UG=Hs.30464 cyclin E2 /FL=gb:AF091433.1 gb:AF102778.1 gb:AF106690.1 gb:NM_004702.1	NM_004702		NP_477097
205038_at	0.001116	zinc finger protein, subfamily 1A, 1 (Ikaros)	BG540504	Hs.54452	NP_006051
205040_at	0.001713	gb:NM_000607.1 /DEF=Homo sapiens orosomucoid 1 (ORM1), mRNA. /FEA=mRNA /GEN=ORM1 /PROD=orosomucoid 1 precursor /DB_XREF=gi:9257231 /UG=Hs.572 orosomucoid 1 /FL=gb:M13692.1 gb:NM_000607.1	NM_000607		NP_000598
205041_s_at	0.007258	gb:NM_000607.1 /DEF=Homo sapiens orosomucoid 1 (ORM1), mRNA. /FEA=mRNA /GEN=ORM1 /PROD=orosomucoid 1 precursor /DB_XREF=gi:9257231 /UG=Hs.572 orosomucoid 1 /FL=gb:M13692.1 gb:NM_000607.1	NM_000607		NP_000598

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205052_at	0.009179	gb:NM_001698.1 /DEF=Homo sapiens AU RNA-binding protein enoyl-Coenzyme A hydratase (AUH), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=AUH /PROD=AU RNA-binding protein enoyl-Coenzyme A hydratase precursor /DB_XREF=gi:4502326 /UG=Hs.81886 AU RNA-binding protein enoyl-Coenzyme A hydratase /FL=gb:NM_001698.1	NM_001698		NP_001689
205070_at	1.26E-04	gb:NM_019071.1 /DEF=Homo sapiens inhibitor of growth family, member 3 (ING3), mRNA. /FEA=mRNA /GEN=ING3 /PROD=inhibitor of growth family, member 3 /DB_XREF=gi:9506658 /UG=Hs.143198 inhibitor of growth family, member 3 /FL=gb:AF074968.1 gb:AY007790.1 gb:NM_019071.1	NM_019071		NP_061944
205072_s_at	0.007943	gb:NM_022406.1 /DEF=Homo sapiens X-ray repair complementing defective repair in Chinese hamster cells 4 (XRCC4), transcript variant 2, mRNA. /FEA=mRNA /GEN=XRCC4 /PROD=X-ray repair cross complementing protein 4, isoform 2 /DB_XREF=gi:12408646 /UG=Hs.150930 X-ray repair complementing defective repair in Chinese hamster cells 4 /FL=gb:U40622.1 gb:AB017445.1 gb:NM_003401.2 gb:NM_022406.1	NM_022406		NP_072044

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205078_at	0.005852	gb:NM_002643.1 /DEF=Homo sapiens phosphatidylinositol glycan, class F (PIGF), mRNA. /FEA=mRNA /GEN=PIGF /PROD=phosphatidylinositol glycan, class F /DB_XREF=gi:4505796 /UG=Hs.166982 phosphatidylinositol glycan, class F /FL=gb:D13435.1 gb:NM_002643.1	NM_002643		NP_775097
205080_at	0.021312	gb:NM_000965.1 /DEF=Homo sapiens retinoic acid receptor, beta (RARβ), mRNA. /FEA=mRNA /GEN=RARB /PROD=retinoic acid receptor, beta /DB_XREF=gi:4506420 /UG=Hs.171495 retinoic acid receptor, beta /FL=gb:NM_000965.1	NM_000965		NP_057236
205098_at	0.016934	chemokine (C-C motif) receptor 1	AI421071	Hs.301921	NP_001286
205105_at	0.041055	gb:NM_002372.1 /DEF=Homo sapiens mannosidase, alpha, class 2A, member 1 (MAN2A1), mRNA. /FEA=mRNA /GEN=MAN2A1 /PROD=mannosidase, alpha, class 2A, member 1 /DB_XREF=gi:4758697 /UG=Hs.32965 mannosidase, alpha, class 2A, member 1 /FL=gb:U31520.1 gb:NM_002372.1 gb:D63998.1	NM_002372		NP_002363
205114_s_at	0.041055	gb:NM_002983.1 /DEF=Homo sapiens small inducible cytokine A3 (homologous to mouse Mip-1a) (SCYA3), mRNA. /FEA=mRNA /GEN=SCYA3 /PROD=small inducible cytokine A3 (homologous to mouse Mip-1a) /DB_XREF=gi:4506842 /UG=Hs.73817 small inducible cytokine A3 (homologous to mouse Mip-1a) /FL=gb:M23452.1 gb:D00044.1 gb:NM_002983.1 gb:M25315.1	NM_002983		NP_002974

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205158_at	0.004733	gb:NM_002937.1 /DEF=Homo sapiens ribonuclease, RNase A family, 4 (RNASE4), mRNA. /FEA=mRNA /GEN=RNASE4 /PROD=ribonuclease, RNase A family, 4 /DB_XREF=gi:4506556 /UG=Hs.283749 ribonuclease, RNase A family, 4 /FL=gb:NM_002937.1 gb:D37931.1	NM_002937		NP_002928
205159_at	0.026891	colony stimulating factor 2 receptor, beta, low-affinity (granulocyte-macrophage)	AV756141	Hs.285401	NP_000386
205173_x_at	1.26E-04	gb:NM_001779.1 /DEF=Homo sapiens CD58 antigen, (lymphocyte function-associated antigen 3) (CD58), mRNA. /FEA=mRNA /GEN=CD58 /PROD=CD58 antigen, (lymphocyte function-associated antigen 3) /DB_XREF=gi:4502676 /UG=Hs.75626 CD58 antigen, (lymphocyte function-associated antigen 3) /FL=gb:NM_001779.1	NM_001779		NP_001770
205174_s_at	0.00161	gb:NM_012413.2 /DEF=Homo sapiens glutaminyl-peptide cyclotransferase (glutaminyl cyclase) (QPCT), mRNA. /FEA=mRNA /GEN=QPCT /PROD=glutaminyl-peptide cyclotransferase precursor /DB_XREF=gi:9257235 /UG=Hs.79033 glutaminyl-peptide cyclotransferase (glutaminyl cyclase) /FL=gb:NM_012413.2	NM_012413		NP_036545
205178_s_at	0.018693	gb:NM_006910.1 /DEF=Homo sapiens retinoblastoma-binding protein 6 (RBBP6), mRNA. /FEA=mRNA /GEN=RBBP6 /PROD=retinoblastoma-binding protein 6 /DB_XREF=gi:5902043 /UG=Hs.85273 retinoblastoma-binding protein 6 /FL=gb:NM_006910.1	NM_006910		NP_008841

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205187_at	0.042466	gb:AF010601.1 /DEF=Homo sapiens SMAD5 (Smad5) mRNA, complete cds. /FEA=mRNA /GEN=Smad5 /PROD=SMAD5 /DB_XREF=gi:2317785 /UG=Hs.37501 MAD (mothers against decapentaplegic, Drosophila) homolog 5 /FL=gb:U59913.1 gb:U73825.1 gb:AF009678.1 gb:AF010601.1 gb:NM_005903.1	AF010601		NP_005894
205191_at	1.26E-04	gb:NM_006915.1 /DEF=Homo sapiens retinitis pigmentosa 2 (X-linked recessive) (RP2), mRNA. /FEA=mRNA /GEN=RP2 /PROD=XRP2 protein /DB_XREF=gi:5902059 /UG=Hs.44766 retinitis pigmentosa 2 (X-linked recessive) /FL=gb:NM_006915.1	NM_006915		NP_008846
205197_s_at	0.011377	ATPase, Cu++ transporting, alpha polypeptide (Menkes syndrome)	BE567813	Hs.606	NP_000043
205202_at	1.26E-04	gb:NM_005389.1 /DEF=Homo sapiens protein-L-isoaspartate (D-aspartate) O-methyltransferase (PCMT1), mRNA. /FEA=mRNA /GEN=PCMT1 /PROD=protein-L-isoaspartate (D-aspartate)O-methyltransferase /DB_XREF=gi:4885538 /UG=Hs.79137 protein-L-isoaspartate (D-aspartate) O-methyltransferase /FL=gb:D25545.1 gb:NM_005389.1	NM_005389		NP_005380
205207_at	0.041795	gb:NM_000600.1 /DEF=Homo sapiens interleukin 6 (interferon, beta 2) (IL6), mRNA. /FEA=mRNA /GEN=IL6 /PROD=interleukin 6 (interferon, beta 2) /DB_XREF=gi:10834983 /UG=Hs.93913 interleukin 6 (interferon, beta 2) /FL=gb:NM_000600.1 gb:M14584.1 gb:M18403.1 gb:M29150.1 gb:M54894.1	NM_000600		NP_000591

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205212_s_at	0.035611	gb:NM_014716.1 /DEF=Homo sapiens KIAA0050 gene product (ACAP1), mRNA. /FEA=mRNA /GEN=ACAP1 /PROD=centaurin beta1 /DB_XREF=gi:7661879 /UG=Hs.108947 KIAA0050 gene product /FL=gb:D30758.1 gb:NM_014716.1	NM_014716		NP_055531
205220_at	0.013223	gb:NM_006018.1 /DEF=Homo sapiens putative chemokine receptor; GTP-binding protein (HM74), mRNA. /FEA=mRNA /GEN=HM74 /PROD=putative chemokine receptor; GTP-bindingprotein /DB_XREF=gi:5174460 /UG=Hs.137555 putative chemokine receptor; GTP-binding protein /FL=gb:NM_006018.1	NM_006018		NP_006009
205230_at	0.006096	gb:NM_014954.1 /DEF=Homo sapiens KIAA0985 protein (KIAA0985), mRNA. /FEA=mRNA /GEN=KIAA0985 /PROD=KIAA0985 protein /DB_XREF=gi:7662431 /UG=Hs.21239 KIAA0985 protein /FL=gb:AB023202.1 gb:NM_014954.1	NM_014954		NP_055769
205245_at	0.013575	gb:NM_016948.1 /DEF=Homo sapiens Tax interaction protein 40 (TAX40), mRNA. /FEA=mRNA /GEN=TAX40 /PROD=Tax interaction protein 40 /DB_XREF=gi:8394416 /UG=Hs.112933 Tax interaction protein 40 /FL=gb:AB041642.1 gb:AF252292.1 gb:AB043634.1 gb:NM_016948.1 gb:AF265565.1	NM_016948		NP_058644
205248_at	1.26E-04	gb:NM_005128.1 /DEF=Homo sapiens chromosome 21 open reading frame 5 (C21ORF5), mRNA. /FEA=mRNA /GEN=C21ORF5 /PROD=chromosome 21 open reading frame 5 /DB_XREF=gi:4826652 /UG=Hs.129781 chromosome 21 open reading frame 5 /FL=gb:NM_005128.1	NM_005128		NP_005119

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205251_at	1.26E-04	gb:NM_022817.1 /DEF=Homo sapiens period (Drosophila) homolog 2 (PER2), transcript variant 1, mRNA. /FEA=mRNA /GEN=PER2 /PROD=period 2, isoform 2 /DB_XREF=gi:12707561 /UG=Hs.153405 period (Drosophila) homolog 2 /FL=gb:NM_022817.1	NM_022817		NP_073728
205255_x_at	7.58E-04	gb:NM_003202.1 /DEF=Homo sapiens transcription factor 7 (T-cell specific, HMG-box) (TCF7), mRNA. /FEA=mRNA /GEN=TCF7 /PROD=transcription factor 7 (T-cell specific,HMG-box) /DB_XREF=gi:4507402 /UG=Hs.169294 transcription factor 7 (T-cell specific, HMG-box) /FL=gb:NM_003202.1	NM_003202		NP_003193
205263_at	7.58E-04	gb:AF082283.1 /DEF=Homo sapiens CARD-containing apoptotic signaling protein (BCL10) mRNA, complete cds. /FEA=mRNA /GEN=BCL10 /PROD=CARD-containing apoptotic signaling protein /DB_XREF=gi:4092066 /UG=Hs.193516 B-cell CLLlymphoma 10 /FL=gb:AF082283.1 gb:AF057700.1 gb:AF100338.1 gb:NM_003921.1 gb:AF127386.1 gb:AF134395.1 gb:AF105066.1	AF082283		NP_003912
205267_at	0.030771	gb:NM_006235.1 /DEF=Homo sapiens POU domain, class 2, associating factor 1 (POU2AF1), mRNA. /FEA=mRNA /GEN=POU2AF1 /PROD=POU domain, class 2, associating factor 1 /DB_XREF=gi:5453933 /UG=Hs.2407 POU domain, class 2, associating factor 1 /FL=gb:NM_006235.1	NM_006235		NP_006226
205269_at	0.010205	lymphocyte cytosolic protein 2 (SH2 domain containing leukocyte protein of 76kDa)	AI123251	Hs.2488	NP_005556

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205270_s_at	0.02145	gb:NM_005565.2 /DEF=Homo sapiens lymphocyte cytosolic protein 2 (SH2 domain-containing leukocyte protein of 76kD) (LCP2), mRNA. /FEA=mRNA /GEN=LCP2 /PROD=lymphocyte cytosolic protein 2 /DB_XREF=gi:7382491 /UG=Hs.2488 lymphocyte cytosolic protein 2 (SH2 domain-containing leukocyte protein of 76kD) /FL=gb:NM_005565.2 gb:U20158.1	NM_005565		NP_005556
205273_s_at	0.017832	gb:NM_014968.1 /DEF=Homo sapiens KIAA1104 protein (KIAA1104), mRNA. /FEA=mRNA /GEN=KIAA1104 /PROD=KIAA1104 protein /DB_XREF=gi:7662487 /UG=Hs.279039 KIAA1104 protein /FL=gb:AB029027.1 gb:NM_014968.1	NM_014968		NP_055783
205281_s_at	0.02145	gb:NM_002641.1 /DEF=Homo sapiens phosphatidylinositol glycan, class A (paroxysmal nocturnal hemoglobinuria) (PIGA), transcript variant 1, mRNA. /FEA=mRNA /GEN=PIGA /PROD=phosphatidylinositol glycan, class A isoform 1 /DB_XREF=gi:11863129 /UG=Hs.51 phosphatidylinositol glycan, class A (paroxysmal nocturnal hemoglobinuria) /FL=gb:NM_002641.1 gb:D11466.1	NM_002641		NP_065206
205285_s_at	0.013088	FYN binding protein (FYB-120/130)	AI633888	Hs.58435	NP_001456

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
205292_s_at	0.026891	gb:NM_002137.1 /DEF=Homo sapiens heterogeneous nuclear ribonucleoprotein A2B1 (HNRPA2B1), mRNA. /FEA=mRNA /GEN=HNRPA2B1 /PROD=heterogeneous nuclear ribonucleoprotein A2B1 /DB_XREF=gi:4504446 /UG=Hs.75598 heterogeneous nuclear ribonucleoprotein A2B1 /FL=gb:NM_002137.1	NM_002137		NP_112533
205297_s_at	6.68E-04	gb:NM_000626.1 /DEF=Homo sapiens CD79B antigen (immunoglobulin-associated beta) (CD79B), transcript variant 1, mRNA. /FEA=mRNA /GEN=CD79B /PROD=CD79B antigen, isoform 1 precursor /DB_XREF=gi:11038673 /UG=Hs.89575 CD79B antigen (immunoglobulin-associated beta) /FL=gb:NM_000626.1 gb:M80461.1 gb:M89957.1	NM_000626		NP_067613
205300_s_at	0.005852	gb:NM_022717.1 /DEF=Homo sapiens U1-snRNP binding protein homolog (70kD) (U1SNRNPBP), transcript variant 2, mRNA. /FEA=mRNA /GEN=U1SNRNPBP /PROD=U1-snRNP binding protein homolog (70kD), isoforma /DB_XREF=gi:13027641 /UG=Hs.93502 U1-snRNP binding protein homolog (70kD) /FL=gb:U44799.1 gb:NM_022717.1	NM_022717		NP_851034
205306_x_at	0.007258	opsin 3 (encephalopsin, panopsin)	AI074145	Hs.107318	NP_003670
205322_s_at	0.037567	metal-regulatory transcription factor 1	AW182367	Hs.211581	NP_005946
205323_s_at	0.041055	gb:NM_005955.1 /DEF=Homo sapiens metal-regulatory transcription factor 1 (MTF1), mRNA. /FEA=mRNA /GEN=MTF1 /PROD=metal-regulatory transcription factor 1 /DB_XREF=gi:5174588 /UG=Hs.211581 metal-regulatory transcription factor 1 /FL=gb:NM_005955.1	NM_005955		NP_005946

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205349_at	3.26E-04	gb:NM_002068.1 /DEF=Homo sapiens guanine nucleotide binding protein (G protein), alpha 15 (Gq class) (GNA15), mRNA. /FEA=mRNA /GEN=GNA15 /PROD=guanine nucleotide binding protein (G protein),alpha 15 (Gq class) /DB_XREF=gi:4504038 /UG=Hs.73797 guanine nucleotide binding protein (G protein), alpha 15 (Gq class) /FL=gb:M63904.1 gb:NM_002068.1	NM_002068		NP_002059
205370_x_at	0.001116	gb:NM_001918.1 /DEF=Homo sapiens dihydrolipoamide branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex; maple syrup urine disease) (DBT), mRNA. /FEA=mRNA /GEN=DBT /PROD=dihydrolipoamide branched chain transacylase (E2component of branched chain keto acid dehydrogenasecomplex; maple syrup urine disease) /DB_XREF=gi:4503264 /UG=Hs.139410 dihydrolipoamide branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex; maple syrup urine disease) /FL=gb:J03208.1 gb:NM_001918.1 gb:M27093.1	NM_001918		NP_001909
205383_s_at	0.004341	gb:NM_015642.1 /DEF=Homo sapiens zinc finger protein 288 (ZNF288), mRNA. /FEA=mRNA /GEN=ZNF288 /PROD=zinc finger protein 288 /DB_XREF=gi:7661651 /UG=Hs.159456 zinc finger protein 288 /FL=gb:AL050276.1 gb:NM_015642.1	NM_015642		NP_056457

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205394_at	0.00545	gb:NM_001274.1 /DEF=Homo sapiens CHK1 (checkpoint, S.pombe) homolog (CHEK1), mRNA. /FEA=mRNA /GEN=CHK1 /PROD=CHK1 (checkpoint, S.pombe) homolog /DB_XREF=gi:4502802 /UG=Hs.20295 CHK1 (checkpoint, S.pombe) homolog /FL=gb:AF016582.1 gb:NM_001274.1	NM_001274		NP_001265
205396_at	0.021448	DKFZP586N0721 protein	BF971416	Hs.99843	NP_056215
205401_at	0.003171	gb:NM_003659.1 /DEF=Homo sapiens alkylglycerone phosphate synthase (AGPS), mRNA. /FEA=mRNA /GEN=AGPS /PROD=alkylglycerone phosphate synthase precursor /DB_XREF=gi:4501992 /UG=Hs.22580 alkylglycerone phosphate synthase /FL=gb:NM_003659.1	NM_003659		NP_003650
205403_at	3.26E-04	gb:NM_004633.1 /DEF=Homo sapiens interleukin 1 receptor, type II (IL1R2), mRNA. /FEA=mRNA /GEN=IL1R2 /PROD=interleukin 1 receptor, type II /DB_XREF=gi:4758597 /UG=Hs.25333 interleukin 1 receptor, type II /FL=gb:U74649.1 gb:NM_004633.1	NM_004633		NP_775465
205418_at	0.003522	gb:NM_002005.2 /DEF=Homo sapiens feline sarcoma (Snyder-Theilen) viral (v-fes)Fujinami avian sarcoma (PRCII) viral (v-fps) oncogene homolog (FES), mRNA. /FEA=mRNA /GEN=FES /PROD=V-FES feline sarcoma viralV-FPS fujinami aviansarcoma viral oncogene homolog /DB_XREF=gi:13376997 /UG=Hs.7636 feline sarcoma (Snyder-Theilen) viral (v-fes)Fujinami avian sarcoma (PRCII) viral (v-fps) oncogene homolog /FL=gb:NM_002005.2	NM_002005		NP_001996

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205423_at	1.26E-04	gb:NM_001127.1 /DEF=Homo sapiens adaptor-related protein complex 1, beta 1 subunit (AP1B1), mRNA. /FEA=mRNA /GEN=AP1B1 /PROD=adaptor-related protein complex 1, beta 1 subunit /DB_XREF=gi:4501972 /UG=Hs.89576 adaptor-related protein complex 1, beta 1 subunit /FL=gb:L13939.1 gb:NM_001127.1	NM_001127		NP_663782
205428_s_at	0.027633	gb:NM_001740.2 /DEF=Homo sapiens calbindin 2, (29kD, calretinin) (CALB2), transcript variant CALB2, mRNA. /FEA=mRNA /GEN=CALB2 /PROD=calbindin 2, full length protein isoform /DB_XREF=gi:6031158 /UG=Hs.106857 calbindin 2, (29kD, calretinin) /FL=gb:NM_001740.2	NM_001740		NP_009019
205436_s_at	0.003529	gb:NM_002105.1 /DEF=Homo sapiens H2A histone family, member X (H2AFX), mRNA. /FEA=mRNA /GEN=H2AFX /PROD=H2A histone family, member X /DB_XREF=gi:4504252 /UG=Hs.147097 H2A histone family, member X /FL=gb:BC004915.1 gb:NM_002105.1	NM_002105		NP_002096
205444_at	0.041795	gb:NM_004320.1 /DEF=Homo sapiens ATPase, Ca++ transporting, cardiac muscle, fast twitch 1 (ATP2A1), mRNA. /FEA=mRNA /GEN=ATP2A1 /PROD=ATPase, Ca++ transporting, cardiac muscle, fasttwitch 1 /DB_XREF=gi:10835219 /UG=Hs.183075 ATPase, Ca++ transporting, cardiac muscle, fast twitch 1 /FL=gb:NM_004320.1	NM_004320		NP_775293

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205451_at	0.013575	gb:NM_005938.1 /DEF=Homo sapiens myeloidlymphoid or mixed-lineage leukemia (trithorax (Drosophila) homolog); translocated to, 7 (MLLT7), mRNA. /FEA=mRNA /GEN=MLLT7 /PROD=myeloidlymphoid or mixed-lineage leukemia(trithorax (Drosophila) homolog); translocated to, 7 /DB_XREF=gi:5174578 /UG=Hs.239663 myeloidlymphoid or mixed-lineage leukemia (trithorax (Drosophila) homolog); translocated to, 7 /FL=gb:NM_005938.1	NM_005938		NP_005929
205456_at	0.016934	gb:NM_000733.1 /DEF=Homo sapiens CD3E antigen, epsilon polypeptide (TiT3 complex) (CD3E), mRNA. /FEA=mRNA /GEN=CD3E /PROD=CD3E antigen, epsilon polypeptide (TiT3complex) /DB_XREF=gi:4502670 /UG=Hs.3003 CD3E antigen, epsilon polypeptide (TiT3 complex) /FL=gb:NM_000733.1	NM_000733		NP_000724
205462_s_at	0.022795	gb:NM_002149.1 /DEF=Homo sapiens hippocalcin-like 1 (HPCAL1), mRNA. /FEA=mRNA /GEN=HPCAL1 /PROD=hippocalcin-like 1 /DB_XREF=gi:4504474 /UG=Hs.3618 hippocalcin-like 1 /FL=gb:NM_002149.1 gb:D16227.1	NM_002149		NP_602293
205467_at	0.007779	gb:NM_001230.1 /DEF=Homo sapiens caspase 10, apoptosis-related cysteine protease (CASP10), mRNA. /FEA=mRNA /GEN=CASP10 /PROD=caspase 10, apoptosis-related cysteine protease /DB_XREF=gi:4502568 /UG=Hs.5353 caspase 10, apoptosis-related cysteine protease /FL=gb:U60519.1 gb:NM_001230.1	NM_001230		NP_116759

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205469_s_at	0.005852	gb:NM_002200.1 /DEF=Homo sapiens interferon regulatory factor 5 (IRF5), mRNA. /FEA=mRNA /GEN=IRF5 /PROD=interferon regulatory factor 5 /DB_XREF=gi:4504726 /UG=Hs.54434 interferon regulatory factor 5 /FL=gb:U51127.1 gb:NM_002200.1	NM_002200		NP_116032
205471_s_at	0.027633	dachshund homolog (Drosophila)	AW772082	Hs.63931	NP_542938
205480_s_at	1.26E-04	gb:NM_006759.2 /DEF=Homo sapiens UDP-glucose pyrophosphorylase 2 (UGP2), mRNA. /FEA=mRNA /GEN=UGP2 /PROD=UDP-glucose pyrophosphorylase 2 /DB_XREF=gi:13027637 /UG=Hs.77837 UDP-glucose pyrophosphorylase 2 /FL=gb:NM_006759.2	NM_006759		NP_006750
205495_s_at	0.003171	gb:NM_006433.2 /DEF=Homo sapiens granulysin (GNLY), transcript variant NKG5, mRNA. /FEA=mRNA /GEN=GNLY /PROD=granulysin, isoform NKG5 /DB_XREF=gi:7108343 /UG=Hs.105806 granulysin /FL=gb:NM_006433.2	NM_006433		NP_036615
205504_at	7.58E-04	gb:NM_000061.1 /DEF=Homo sapiens Bruton agammaglobulinemia tyrosine kinase (BTK), mRNA. /FEA=mRNA /GEN=BTK /PROD=Bruton agammaglobulinemia tyrosine kinase /DB_XREF=gi:4557376 /UG=Hs.159494 Bruton agammaglobulinemia tyrosine kinase /FL=gb:NM_000061.1 gb:AF153756.1 gb:AF153757.1 gb:AF153758.1 gb:AF153759.1 gb:AF153760.1 gb:AF153761.1 gb:AF153762.1 gb:AF153364.1	NM_000061		NP_000052

Gene Identifier	p-value	D scription	Gen Accession No.	Unigene Accession No.	Protein Accession No.
205510_s_at	0.00161	gb:NM_017976.1 /DEF=Homo sapiens hypothetical protein FLJ10038 (FLJ10038), mRNA. /FEA=mRNA /GEN=FLJ10038 /PROD=hypothetical protein FLJ10038 /DB_XREF=gi:8922197 /UG=Hs.181202 hypothetical protein FLJ10038 /FL=gb:NM_017976.1	NM_017976		NP_060446
205513_at	5.03E-04	gb:NM_001062.1 /DEF=Homo sapiens transcobalamin I (vitamin B12 binding protein, R binder family) (TCN1), mRNA. /FEA=mRNA /GEN=TCN1 /PROD=transcobalamin I (vitamin B12 binding protein, Rbinder family) /DB_XREF=gi:4507406 /UG=Hs.2012 transcobalamin I (vitamin B12 binding protein, R binder family) /FL=gb:J05068.1 gb:NM_001062.1	NM_001062		NP_001053
205525_at	0.017832	gb:NM_018495.3 /DEF=Homo sapiens NAG22 protein (LOC55873), mRNA. /FEA=mRNA /GEN=LOC55873 /PROD=NAG22 protein /DB_XREF=gi:13236500 /UG=Hs.283080 NAG22 protein /FL=gb:AF247820.3 gb:NM_018495.3	NM_018495		
205539_at	0.004341	gb:NM_006576.1 /DEF=Homo sapiens advillin (AVIL), mRNA. /FEA=mRNA /GEN=AVIL /PROD=advillin /DB_XREF=gi:5729735 /UG=Hs.47344 advillin /FL=gb:AF041449.1 gb:NM_006576.1	NM_006576		NP_006567

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205550_s_at	2.05E-04	gb:NM_004899.1 /DEF=Homo sapiens brain and reproductive organ-expressed (TNFRSF1A modulator) (BRE), mRNA. /FEA=mRNA /GEN=BRE /PROD=brain and reproductive organ-expressed (TNFRSF1A modulator) /DB_XREF=gi:4757871 /UG=Hs.80426 brain and reproductive organ-expressed (TNFRSF1A modulator) /FL=gb:BC001251.1 gb:NM_004899.1 gb:L38616.1	NM_004899		NP_004890
205557_at	3.09E-04	gb:NM_001725.1 /DEF=Homo sapiens bactericidal permeability-increasing protein (BPI), mRNA. /FEA=mRNA /GEN=BPI /PROD=bactericidal permeability-increasing protein precursor /DB_XREF=gi:4502446 /UG=Hs.89535 bactericidal permeability-increasing protein /FL=gb:AF322588.1 gb:J04739.1 gb:NM_001725.1	NM_001725		NP_001716
205568_at	0.033381	gb:NM_020980.2 /DEF=Homo sapiens aquaporin 9 (AQP9), mRNA. /FEA=mRNA /GEN=AQP9 /PROD=aquaporin 9 /DB_XREF=gi:11038652 /UG=Hs.104624 aquaporin 9 /FL=gb:NM_020980.2 gb:AB008775.1 gb:AF016495.1	NM_020980		NP_066190
205570_at	0.026891	gb:NM_005028.2 /DEF=Homo sapiens phosphatidylinositol-4-phosphate 5-kinase, type II, alpha (PIP5K2A), mRNA. /FEA=mRNA /GEN=PIP5K2A /PROD=phosphatidylinositol-4-phosphate 5-kinase, type II, alpha /DB_XREF=gi:6857819 /UG=Hs.108966 phosphatidylinositol-4-phosphate 5-kinase, type II, alpha /FL=gb:U14957.2 gb:NM_005028.2	NM_005028		NP_005019

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205575_at	5.03E-04	gb:NM_006688.1 /DEF=Homo sapiens C1q-related factor (CRF), mRNA. /FEA=mRNA /GEN=CRF /PROD=C1q-related factor /DB_XREF=gi:5729784 /UG=Hs.134012 C1q-related factor /FL=gb:AF095154.1 gb:NM_006688.1	NM_006688		NP_006679
205588_s_at	0.011862	gb:NM_007045.1 /DEF=Homo sapiens FGFR1 oncogene partner (FOP), mRNA. /FEA=mRNA /GEN=FOP /PROD=FGFR1 oncogene partner /DB_XREF=gi:5901953 /UG=Hs.180296 FGFR1 oncogene partner /FL=gb:NM_007045.1	NM_007045		NP_008976
205590_at	7.58E-04	gb:NM_005739.2 /DEF=Homo sapiens RAS guanyl releasing protein 1 (calcium and DAG-regulated) (RASGRP1), mRNA. /FEA=mRNA /GEN=RASGRP1 /PROD=RAS guanyl releasing protein 1 /DB_XREF=gi:6382080 /UG=Hs.182591 RAS guanyl releasing protein 1 (calcium and DAG-regulated) /FL=gb:AF081195.1 gb:AF106071.1 gb:NM_005739.2	NM_005739		NP_005730
205594_at	0.041055	gb:NM_014897.1 /DEF=Homo sapiens KIAA0924 protein (KIAA0924), mRNA. /FEA=mRNA /GEN=KIAA0924 /PROD=KIAA0924 protein /DB_XREF=gi:7662383 /UG=Hs.190386 KIAA0924 protein /FL=gb:AB023141.1 gb:NM_014897.1	NM_014897		NP_055712
205596_s_at	0.010205	gb:AY014180.1 /DEF=Homo sapiens E3 ubiquitin ligase Smurf2 mRNA, complete cds. /FEA=mRNA /PROD=E3 ubiquitin ligase Smurf2 /DB_XREF=gi:12408118 /UG=Hs.194477 E3 ubiquitin ligase SMURF2 /FL=gb:AF301463.1 gb:AF310676.1 gb:NM_022739.1 gb:AY014180.1	AY014180		NP_073576

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205603_s_at	0.00161	gb:NM_007309.1 /DEF=Homo sapiens diaphanous (Drosophila, homolog) 2 (DIAPH2), transcript variant 12C, mRNA. /FEA=mRNA /GEN=DIAPH2 /PROD=diaphanous 2 isoform 12C /DB_XREF=gi:6382070 /UG=Hs.226483 diaphanous (Drosophila, homolog) 2 /FL=gb:NM_007309.1	NM_007309		NP_009293
205607_s_at	5.03E-04	gb:NM_020423.1 /DEF=Homo sapiens hypothetical protein LOC57147 (LOC57147), mRNA. /FEA=mRNA /GEN=LOC57147 /PROD=hypothetical protein LOC57147 /DB_XREF=gi:9967093 /UG=Hs.24243 hypothetical protein LOC57147 /FL=gb:NM_020423.1	NM_020423		NP_065156
205627_at	0.026891	gb:NM_001785.1 /DEF=Homo sapiens cytidine deaminase (CDA), mRNA. /FEA=mRNA /GEN=CDA /PROD=cytidine deaminase /DB_XREF=gi:11386156 /UG=Hs.72924 cytidine deaminase /FL=gb:NM_001785.1 gb:L27943.1	NM_001785		NP_001776
205632_s_at	0.003171	gb:NM_003558.1 /DEF=Homo sapiens phosphatidylinositol-4-phosphate 5-kinase, type I, beta (PIP5K1B), mRNA. /FEA=mRNA /GEN=PIP5K1B /PROD=phosphatidylinositol-4-phosphate 5-kinase, type I, beta /DB_XREF=gi:4505816 /UG=Hs.78406 phosphatidylinositol 4-phosphate 5-kinase, type I, beta /FL=gb:NM_003558.1	NM_003558		NP_003549

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205633_s_at	1.26E-04	gb:NM_000688.1 /DEF=Homo sapiens aminolevulinate, delta-, synthase 1 (ALAS1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=ALAS1 /PROD=aminolevulinate, delta-, synthase 1 /DB_XREF=gi:4502024 /UG=Hs.78712 aminolevulinate, delta-, synthase 1 /FL=gb:NM_000688.1	NM_000688		NP_000679
205636_at	0.037567	Consensus includes gb:AF036269.1 /DEF=Homo sapiens EEN-B2-L1 mRNA, complete cds. /FEA=mRNA /GEN=EEN-B2-L1 /PROD=EEN-B2-L1 /DB_XREF=gi:2921409 /UG=Hs.80315 SH3-domain GRB2-like 3 /FL=gb:AF036269.1 gb:AF036270.1 gb:AF036271.1 gb:NM_003027.1	AF036269		NP_003018
205639_at	1.26E-04	gb:NM_001637.1 /DEF=Homo sapiens acyloxyacyl hydrolase (neutrophil) (AOAH), mRNA. /FEA=mRNA /GEN=AOAH /PROD=acyloxyacyl hydrolase precursor /DB_XREF=gi:4502114 /UG=Hs.82542 acyloxyacyl hydrolase (neutrophil) /FL=gb:M62840.1 gb:NM_001637.1	NM_001637		NP_001628
205640_at	0.001323	gb:NM_000694.1 /DEF=Homo sapiens aldehyde dehydrogenase 3 family, member B1 (ALDH3B1), mRNA. /FEA=mRNA /GEN=ALDH3B1 /PROD=aldehyde dehydrogenase 3B1 /DB_XREF=gi:4502042 /UG=Hs.83155 aldehyde dehydrogenase 3 family, member B1 /FL=gb:NM_000694.1 gb:U10868.1	NM_000694		NP_000685

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205645_at	0.021448	gb:NM_004726.1 /DEF=Homo sapiens RALBP1 associated Eps domain containing 2 (REPS2), mRNA. /FEA=mRNA /GEN=REPS2 /PROD=RALBP1 associated Eps domain containing 2 /DB_XREF=gi:4758943 /UG=Hs.80667 RALBP1 associated Eps domain containing 2 /FL=gb:AF010233.1 gb:NM_004726.1	NM_004726		NP_004717
205653_at	0.041795	gb:NM_001911.1 /DEF=Homo sapiens cathepsin G (CTSG), mRNA. /FEA=mRNA /GEN=CTSG /PROD=cathepsin G /DB_XREF=gi:4503148 /UG=Hs.100764 cathepsin G /FL=gb:M16117.1 gb:NM_001911.1	NM_001911		NP_001902
205681_at	0.033381	gb:NM_004049.1 /DEF=Homo sapiens BCL2-related protein A1 (BCL2A1), mRNA. /FEA=mRNA /GEN=BCL2A1 /PROD=BCL2-related protein A1 /DB_XREF=gi:4757839 /UG=Hs.227817 BCL2-related protein A1 /FL=gb:U27467.1 gb:U29680.1 gb:NM_004049.1	NM_004049		NP_004040
205684_s_at	0.004341	gb:NM_017925.1 /DEF=Homo sapiens hypothetical protein FLJ20686 (FLJ20686), mRNA. /FEA=mRNA /GEN=FLJ20686 /PROD=hypothetical protein FLJ20686 /DB_XREF=gi:8923616 /UG=Hs.271480 hypothetical protein FLJ20686 /FL=gb:NM_017925.1	NM_017925		NP_060395
205690_s_at	0.001116	gb:NM_003910.1 /DEF=Homo sapiens maternal G10 transcript (G10), mRNA. /FEA=mRNA /GEN=G10 /PROD=maternal G10 transcript /DB_XREF=gi:4503836 /UG=Hs.330310 maternal G10 transcript /FL=gb:NM_003910.1 gb:U11861.1	NM_003910		NP_003901

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205698_s_at	0.026891	gb:NM_002758.1 /DEF=Homo sapiens mitogen-activated protein kinase kinase 6 (MAP2K6), mRNA. /FEA=mRNA /GEN=MAP2K6 /PROD=mitogen-activated protein kinase kinase 6 /DB_XREF=gi:10835026 /UG=Hs.118825 mitogen-activated protein kinase kinase 6 /FL=gb:NM_002758.1 gb:U39656.1 gb:U39657.1 gb:U39065.1 gb:U49732.1 gb:D87905.1	NM_002758		NP_114365
205702_at	0.03878	gb:NM_006608.1 /DEF=Homo sapiens putative homeodomain transcription factor (PHTF1), mRNA. /FEA=mRNA /GEN=PHTF1 /PROD=putative homeodomain transcription factor /DB_XREF=gi:5729975 /UG=Hs.123637 putative homeodomain transcription factor /FL=gb:NM_006608.1	NM_006608		NP_006599
205708_s_at	0.004631	ESTs, Highly similar to TRL2_HUMAN Long transient receptor potential channel 2 (LTrpC2) (Transient receptor potential channel 7) (TrpC7) [H.sapiens]	AI051254	Hs.408566	
205711_x_at	0.02145	gb:NM_005174.1 /DEF=Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F1 complex, gamma polypeptide 1 (ATP5C1), mRNA. /FEA=mRNA /GEN=ATP5C1 /PROD=ATP synthase, H ⁺ transporting, mitochondrial F1 complex, gamma polypeptide 1 /DB_XREF=gi:4885078 /UG=Hs.155433 ATP synthase, H ⁺ transporting, mitochondrial F1 complex, gamma polypeptide 1 /FL=gb:D16563.1 gb:NM_005174.1	NM_005174		NP_005165

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205715_at	1.26E-04	gb:NM_004334.1 /DEF=Homo sapiens bone marrow stromal cell antigen 1 (BST1), mRNA. /FEA=mRNA /GEN=BST1 /PROD=bone marrow stromal cell antigen 1 precursor /DB_XREF=gi:4757873 /UG=Hs.169998 bone marrow stromal cell antigen 1 /FL=gb:NM_004334.1 gb:D21878.1	NM_004334		NP_004325
205716_at	0.00228	gb:NM_018843.1 /DEF=Homo sapiens mitochondrial carrier family protein (LOC55972), mRNA. /FEA=mRNA /GEN=LOC55972 /PROD=mitochondrial carrier family protein /DB_XREF=gi:10047121 /UG=Hs.172294 mitochondrial carrier family protein /FL=gb:NM_018843.1 gb:AF125531.1	NM_018843		NP_061331
205726_at	1.26E-04	gb:NM_006729.1 /DEF=Homo sapiens diaphanous (Drosophila, homolog) 2 (DIAPH2), transcript variant 156, mRNA. /FEA=mRNA /GEN=DIAPH2 /PROD=diaphanous 2 isoform 156 /DB_XREF=gi:5803002 /UG=Hs.226483 diaphanous (Drosophila, homolog) 2 /FL=gb:NM_006729.1	NM_006729		
205732_s_at	0.043705	gb:NM_006540.1 /DEF=Homo sapiens nuclear receptor coactivator 2 (NCOA2), mRNA. /FEA=mRNA /GEN=NCOA2 /PROD=nuclear receptor coactivator 2 /DB_XREF=gi:5729857 /UG=Hs.29131 nuclear receptor coactivator 2 /FL=gb:NM_006540.1	NM_006540		NP_006531

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205740_s_at	3.26E-04	gb:NM_024321.1 /DEF=Homo sapiens hypothetical protein MGC10433 (MGC10433), mRNA. /FEA=mRNA /GEN=MGC10433 /PROD=hypothetical protein MGC10433 /DB_XREF=gi:13236562 /UG=Hs.5086 hypothetical protein MGC10433 /FL=gb:BC002868.1 gb:NM_024321.1	NM_024321		NP_077297
205756_s_at	0.00228	gb:NM_000132.2 /DEF=Homo sapiens coagulation factor VIII, procoagulant component (hemophilia A) (F8), transcript variant 1, mRNA. /FEA=mRNA /GEN=F8 /PROD=coagulation factor VIII, procoagulant component, isoform a precursor /DB_XREF=gi:10518504 /UG=Hs.79345 coagulation factor VIII, procoagulant component (hemophilia A) /FL=gb:NM_000132.2 gb:M14113.1	NM_000132		NP_063916
205760_s_at	0.008036	gb:NM_016821.1 /DEF=Homo sapiens 8-oxoguanine DNA glycosylase (OGG1), nuclear gene encoding mitochondrial protein, transcript variant 2a, mRNA. /FEA=mRNA /GEN=OGG1 /PROD=8-oxoguanine DNA glycosylase, isoform 2a /DB_XREF=gi:8670533 /UG=Hs.96398 8-oxoguanine DNA glycosylase /FL=gb:U88620.1 gb:NM_016821.1	NM_016821		NP_058438
205773_at	0.006256	gb:NM_014912.1 /DEF=Homo sapiens KIAA0940 protein (KIAA0940), mRNA. /FEA=mRNA /GEN=KIAA0940 /PROD=KIAA0940 protein /DB_XREF=gi:7662391 /UG=Hs.131945 KIAA0940 protein /FL=gb:AB023157.1 gb:NM_014912.1	NM_014912		NP_055727

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205776_at	0.016495	gb:NM_001461.1 /DEF=Homo sapiens flavin containing monooxygenase 5 (FMO5), mRNA. /FEA=mRNA /GEN=FMO5 /PROD=flavin containing monooxygenase 5 /DB_XREF=gi:4503760 /UG=Hs.14286 flavin containing monooxygenase 5 /FL=gb:NM_001461.1 gb:L37080.1	NM_001461		NP_001452
205786_s_at	1.26E-04	gb:NM_000632.2 /DEF=Homo sapiens integrin, alpha M (complement component receptor 3, alpha; also known as CD11b (p170), macrophage antigen alpha polypeptide) (ITGAM), mRNA. /FEA=mRNA /GEN=ITGAM /PROD=integrin alpha M precursor /DB_XREF=gi:6006013 /UG=Hs.172631 integrin, alpha M (complement component receptor 3, alpha; also known as CD11b (p170), macrophage antigen alpha polypeptide) /FL=gb:J03925.1 gb:NM_000632.2	NM_000632		NP_000623
205787_x_at	0.046906	KIAA0663 gene product	AI803216	Hs.17969	NP_055642
205788_s_at	0.013223	gb:NM_014827.1 /DEF=Homo sapiens KIAA0663 gene product (KIAA0663), mRNA. /FEA=mRNA /GEN=KIAA0663 /PROD=KIAA0663 gene product /DB_XREF=gi:7662231 /UG=Hs.17969 KIAA0663 gene product /FL=gb:AB014563.1 gb:NM_014827.1	NM_014827		NP_055642
205790_at	0.041055	gb:NM_003726.1 /DEF=Homo sapiens src kinase-associated phosphoprotein of 55 kDa (SKAP55), mRNA. /FEA=mRNA /GEN=SKAP55 /PROD=src kinase associated phosphoprotein of 55 kDa /DB_XREF=gi:4506964 /UG=Hs.19126 src kinase-associated phosphoprotein of 55 kDa /FL=gb:NM_003726.1	NM_003726		NP_003717

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205791_x_at	0.035773	gb:NM_006300.1 /DEF=Homo sapiens zinc finger protein 230 (ZNF230), mRNA. /FEA=mRNA /GEN=ZNF230 /PROD=zinc finger protein 230 /DB_XREF=gi:5454181 /UG=Hs.193583 zinc finger protein 230 /FL=gb:U95044.1 gb:NM_006300.1	NM_006300		NP_006291
205798_at	1.26E-04	gb:NM_002185.1 /DEF=Homo sapiens interleukin 7 receptor (IL7R), mRNA. /FEA=mRNA /GEN=IL7R /PROD=interleukin 7 receptor /DB_XREF=gi:4504678 /UG=Hs.237868 interleukin 7 receptor /FL=gb:M29696.1 gb:NM_002185.1	NM_002185		NP_002176
205812_s_at	0.010205	gb:NM_006588.1 /DEF=Homo sapiens sulfotransferase family, cytosolic, 1C, member 2 (SULT1C2), mRNA. /FEA=mRNA /GEN=SULT1C2 /PROD=SULT1C sulfotransferase /DB_XREF=gi:5730070 /UG=Hs.312644 sulfotransferase family, cytosolic, 1C, member 2 /FL=gb:AF055584.1 gb:NM_006588.1	NM_006588		NP_006579
205821_at	0.016426	gb:NM_007360.1 /DEF=Homo sapiens DNA segment on chromosome 12 (unique) 2489 expressed sequence (D12S2489E), mRNA. /FEA=mRNA /GEN=D12S2489E /PROD=NKG2-D type II integral membrane protein /DB_XREF=gi:6679051 /UG=Hs.74085 DNA segment on chromosome 12 (unique) 2489 expressed sequence /FL=gb:NM_007360.1 gb:AF260135.1 gb:AF260136.1	NM_007360		NP_031386

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205831_at	0.033381	gb:NM_001767.1 /DEF=Homo sapiens CD2 antigen (p50), sheep red blood cell receptor (CD2), mRNA. /FEA=mRNA /GEN=CD2 /PROD=CD2 antigen (p50), sheep red blood cellreceptor /DB_XREF=gi:4502652 /UG=Hs.89476 CD2 antigen (p50), sheep red blood cell receptor /FL=gb:M16445.1 gb:M14362.1 gb:M16336.1 gb:NM_001767.1	NM_001767		NP_001758
205832_at	0.008237	gb:NM_016352.1 /DEF=Homo sapiens carboxypeptidase A3 (LOC51200), mRNA. /FEA=mRNA /GEN=LOC51200 /PROD=carboxypeptidase A3 /DB_XREF=gi:10047105 /UG=Hs.93764 carboxypeptidase A3 /FL=gb:NM_016352.1 gb:AF095719.1	NM_016352		NP_057436
205841_at	0.005339	gb:NM_004972.2 /DEF=Homo sapiens Janus kinase 2 (a protein tyrosine kinase) (JAK2), mRNA. /FEA=mRNA /GEN=JAK2 /PROD=Janus kinase 2 /DB_XREF=gi:13325062 /UG=Hs.115541 Janus kinase 2 (a protein tyrosine kinase) /FL=gb:NM_004972.2 gb:AF005216.1 gb:AF058925.1 gb:AF001362.1	NM_004972		NP_004963
205842_s_at	5.03E-04	gb:AF001362.1 /DEF=Homo sapiens Jak2 kinase (JAK2) mRNA, complete cds. /FEA=mRNA /GEN=JAK2 /PROD=Jak2 kinase /DB_XREF=gi:3236321 /UG=Hs.115541 Janus kinase 2 (a protein tyrosine kinase) /FL=gb:NM_004972.2 gb:AF005216.1 gb:AF058925.1 gb:AF001362.1	AF001362		NP_004963

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205844_at	1.26E-04	gb:NM_004666.1 /DEF=Homo sapiens vanin 1 (VNN1), mRNA. /FEA=mRNA /GEN=VNN1 /PROD=vanin 1 /DB_XREF=gi:4759311 /UG=Hs.12114 vanin 1 /FL=gb:NM_004666.1 gb:U39664.1	NM_004666		NP_004657
205856_at	0.001078	gb:NM_015865.1 /DEF=Homo sapiens solute carrier family 14 (urea transporter), member 1 (Kidd blood group) (SLC14A1), mRNA. /FEA=mRNA /GEN=SLC14A1 /PROD=RACH1 /DB_XREF=gi:7706676 /UG=Hs.171731 solute carrier family 14 (urea transporter), member 1 (Kidd blood group) /FL=gb:U35735.1 gb:NM_015865.1	NM_015865		NP_056949
205863_at	1.26E-04	gb:NM_005621.1 /DEF=Homo sapiens S100 calcium-binding protein A12 (calgranulin C) (S100A12), mRNA. /FEA=mRNA /GEN=S100A12 /PROD=S100 calcium-binding protein A12 /DB_XREF=gi:5032058 /UG=Hs.19413 S100 calcium-binding protein A12 (calgranulin C) /FL=gb:D83664.1 gb:D49549.1 gb:NM_005621.1	NM_005621		NP_005612
205865_at	0.033381	gb:NM_005224.1 /DEF=Homo sapiens dead ringer (Drosophila)-like 1 (DRIL1), mRNA. /FEA=mRNA /GEN=DRIL1 /PROD=dead ringer-like 1 /DB_XREF=gi:4885192 /UG=Hs.198515 dead ringer (Drosophila)-like 1 /FL=gb:U88047.1 gb:NM_005224.1	NM_005224		NP_005215

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
205871_at	0.002989	gb:BC005379.1 /DEF=Homo sapiens, Similar to plasminogen, clone MGC:12496, mRNA, complete cds. /FEA=mRNA /PROD=Similar to plasminogen /DB_XREF=gi:13529238 /UG=Hs.262869 plasminogen-like /FL=gb:BC005379.1 gb:M93143.2 gb:NM_002665.1	BC005379		NP_002656
205883_at	0.019879	gb:NM_006006.1 /DEF=Homo sapiens zinc finger protein 145 (Kruppel-like, expressed in promyelocytic leukemia) (ZNF145), mRNA. /FEA=mRNA /GEN=ZNF145 /PROD=zinc finger protein 145 (Kruppel-like, expressed in promyelocytic leukemia) /DB_XREF=gi:5174752 /UG=Hs.37096 zinc finger protein 145 (Kruppel-like, expressed in promyelocytic leukemia) /FL=gb:NM_006006.1	NM_006006		NP_005997
205896_at	2.05E-04	gb:NM_003059.1 /DEF=Homo sapiens solute carrier family 22 (organic cation transporter), member 4 (SLC22A4), mRNA. /FEA=mRNA /GEN=SLC22A4 /PROD=solute carrier family 22 (organic cation transporter), member 4 /DB_XREF=gi:4507002 /UG=Hs.77239 solute carrier family 22 (organic cation transporter), member 4 /FL=gb:AB007448.1 gb:NM_003059.1	NM_003059		NP_003050
205930_at	0.007779	gb:NM_005513.1 /DEF=Homo sapiens general transcription factor IIE, polypeptide 1 (alpha subunit, 56kD) (GTF2E1), mRNA. /FEA=mRNA /GEN=GTF2E1 /PROD=general transcription factor IIE, polypeptide 1(alpha subunit, 56kD) /DB_XREF=gi:5031726 /UG=Hs.145381 general transcription factor IIE, polypeptide 1 (alpha subunit, 56kD) /FL=gb:NM_005513.1	NM_005513		NP_005504

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205936_s_at	6.68E-04	gb:NM_002115.1 /DEF=Homo sapiens hexokinase 3 (white cell) (HK3), mRNA. /FEA=mRNA /GEN=HK3 /PROD=hexokinase 3 (white cell) /DB_XREF=gi:4504394 /UG=Hs.159237 hexokinase 3 (white cell) /FL=gb:U51333.1 gb:NM_002115.1	NM_002115		NP_002106
205945_at	0.013223	gb:NM_000565.1 /DEF=Homo sapiens interleukin 6 receptor (IL6R), mRNA. /FEA=mRNA /GEN=IL6R /PROD=interleukin 6 receptor /DB_XREF=gi:4504672 /UG=Hs.193400 interleukin 6 receptor /FL=gb:NM_000565.1	NM_000565		NP_852004
205950_s_at	0.008237	gb:NM_001738.1 /DEF=Homo sapiens carbonic anhydrase I (CA1), mRNA. /FEA=mRNA /GEN=CA1 /PROD=carbonic anhydrase I /DB_XREF=gi:4502516 /UG=Hs.23118 carbonic anhydrase I /FL=gb:M33987.1 gb:NM_001738.1	NM_001738		NP_001729
205963_s_at	0.005039	gb:NM_005147.1 /DEF=Homo sapiens tumorous imaginal discs (Drosophila) homolog (TID1), mRNA. /FEA=mRNA /GEN=TID1 /PROD=tumorous imaginal discs (Drosophila) homolog /DB_XREF=gi:4827025 /UG=Hs.311057 DnaJ (Hsp40) homolog, subfamily A, member 3 /FL=gb:AF061749.1 gb:NM_005147.1	NM_005147		NP_005138
205967_at	0.033381	gb:NM_003542.2 /DEF=Homo sapiens H4 histone family, member G (H4FG), mRNA. /FEA=mRNA /GEN=H4FG /PROD=H4 histone family, member G /DB_XREF=gi:5579465 /UG=Hs.46423 H4 histone family, member G /FL=gb:NM_003542.2	NM_003542		NP_003533

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
205981_s_at	0.02145	gb:NM_001564.1 /DEF=Homo sapiens inhibitor of growth family, member 1-like (ING1L), mRNA. /FEA=mRNA /GEN=ING1L /PROD=inhibitor of growth 1-like /DB_XREF=gi:4504694 /UG=Hs.107153 inhibitor of growth family, member 1-like /FL=gb:AB012853.1 gb:NM_001564.1 gb:AF053537.1	NM_001564		NP_001555
205988_at	0.026891	gb:NM_003874.1 /DEF=Homo sapiens CD84 antigen (leukocyte antigen) (CD84), mRNA. /FEA=mRNA /GEN=CD84 /PROD=CD84 antigen (leukocyte antigen) /DB_XREF=gi:4502686 /UG=Hs.137548 CD84 antigen (leukocyte antigen) /FL=gb:U82988.1 gb:NM_003874.1 gb:AF054815.1	NM_003874		NP_003865
205992_s_at	0.00161	gb:NM_000585.1 /DEF=Homo sapiens interleukin 15 (IL15), mRNA. /FEA=mRNA /GEN=IL15 /PROD=interleukin 15 /DB_XREF=gi:10835152 /UG=Hs.168132 interleukin 15 /FL=gb:NM_000585.1 gb:U14407.1	NM_000585		NP_751915
205996_s_at	6.79E-04	gb:NM_013411.1 /DEF=Homo sapiens adenylate kinase 2 (AK2), nuclear gene encoding mitochondrial protein, transcript variant AK2B, mRNA. /FEA=mRNA /GEN=AK2 /PROD=adenylate kinase 2 isoform b /DB_XREF=gi:7524345 /UG=Hs.171811 adenylate kinase 2 /FL=gb:U54645.1 gb:NM_013411.1	NM_013411		NP_751949
206003_at	0.033381	gb:NM_014645.1 /DEF=Homo sapiens KIAA0635 gene product (KIAA0635), mRNA. /FEA=mRNA /GEN=KIAA0635 /PROD=KIAA0635 gene product /DB_XREF=gi:7662215 /UG=Hs.185091 KIAA0635 gene product /FL=gb:AB014535.1 gb:NM_014645.1	NM_014645		NP_055460

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206016_at	0.022795	gb:NM_014008.1 /DEF=Homo sapiens JM1 protein (JM1), mRNA. /FEA=mRNA /GEN=JM1 /PROD=JM1 protein /DB_XREF=gi:7661843 /UG=Hs.26333 JM1 protein /FL=gb:BC000972.2 gb:AJ005890.1 gb:NM_014008.1	NM_014008		NP_054727
206026_s_at	0.00161	gb:NM_007115.1 /DEF=Homo sapiens tumor necrosis factor, alpha-induced protein 6 (TNFAIP6), mRNA. /FEA=mRNA /GEN=TNFAIP6 /PROD=tumor necrosis factor, alpha-induced protein 6 /DB_XREF=gi:6005905 /UG=Hs.29352 tumor necrosis factor, alpha-induced protein 6 /FL=gb:NM_007115.1	NM_007115		NP_009046
206028_s_at	7.10E-04	gb:NM_006343.1 /DEF=Homo sapiens c-mer proto-oncogene tyrosine kinase (MERTK), mRNA. /FEA=mRNA /GEN=MERTK /PROD=c-mer proto-oncogene tyrosine kinase /DB_XREF=gi:5453737 /UG=Hs.306178 c-mer proto-oncogene tyrosine kinase /FL=gb:U08023.1 gb:NM_006343.1	NM_006343		NP_006334
206036_s_at	0.022795	gb:NM_002908.1 /DEF=Homo sapiens v-rel avian reticuloendotheliosis viral oncogene homolog (REL), mRNA. /FEA=mRNA /GEN=REL /PROD=v-rel avian reticuloendotheliosis viral oncogene homolog /DB_XREF=gi:4506472 /UG=Hs.44313 v-rel avian reticuloendotheliosis viral oncogene homolog /FL=gb:NM_002908.1	NM_002908		NP_002899

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206042_x_at	0.013223	gb:NM_022804.1 /DEF=Homo sapiens SNRPN upstream reading frame (SNURF), transcript variant 2, mRNA. /FEA=mRNA /GEN=SNURF /PROD=SNRPN upstream reading frame protein /DB_XREF=gi:13027654 /UG=Hs.58606 SNRPN upstream reading frame /FL=gb:NM_005678.2 gb:NM_022804.1 gb:AF101044.1	NM_022804		NP_073715
206050_s_at	0.00228	gb:NM_002939.1 /DEF=Homo sapiens ribonucleaseangiogenin inhibitor (RNH), mRNA. /FEA=mRNA /GEN=RNH /PROD=ribonucleaseangiogenin inhibitor /DB_XREF=gi:4506564 /UG=Hs.75108 ribonucleaseangiogenin inhibitor /FL=gb:M36717.1 gb:NM_002939.1	NM_002939		NP_002930
206052_s_at	1.26E-04	gb:NM_006527.1 /DEF=Homo sapiens Hairpin binding protein, histone (HBP), mRNA. /FEA=mRNA /GEN=HBP /PROD=Hairpin binding protein, histone /DB_XREF=gi:5729861 /UG=Hs.75257 stem-loop (histone) binding protein /FL=gb:U75679.1 gb:NM_006527.1	NM_006527		NP_006518
206056_x_at	0.005039	Human gene for sialophorin (CD43).	NM_003123		NP_003114
206059_at	0.002418	gb:NM_003430.1 /DEF=Homo sapiens zinc finger protein 91 (HPF7, HTF10) (ZNF91), mRNA. /FEA=mRNA /GEN=ZNF91 /PROD=zinc finger protein 91 (HPF7, HTF10) /DB_XREF=gi:4508040 /UG=Hs.8597 zinc finger protein 91 (HPF7, HTF10) /FL=gb:L11672.1 gb:NM_003430.1	NM_003430		NP_003421

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206061_s_at	0.00161	gb:NM_030621.1 /DEF=Homo sapiens helicase-moi (KIAA0928), mRNA. /FEA=mRNA /GEN=KIAA0928 /PROD=helicase-moi /DB_XREF=gi:13449288 /UG=Hs.87889 helicase-moi /FL=gb:NM_030621.1 gb:AB028449.1	NM_030621		NP_803187
206063_x_at	0.021876	peptidylprolyl isomerase (cyclophilin)-like 2	BF001594	Hs.93523	NP_680481
206075_s_at	7.44E-04	gb:NM_001895.1 /DEF=Homo sapiens casein kinase 2, alpha 1 polypeptide (CSNK2A1), mRNA. /FEA=mRNA /GEN=CSNK2A1 /PROD=casein kinase 2, alpha 1 polypeptide /DB_XREF=gi:4503094 /UG=Hs.155140 casein kinase 2, alpha 1 polypeptide /FL=gb:M55265.1 gb:NM_001895.1 gb:J02853.1	NM_001895		NP_808228
206090_s_at	0.012818	gb:NM_018662.1 /DEF=Homo sapiens disrupted in schizophrenia 1 (DISC1), mRNA. /FEA=mRNA /GEN=DISC1 /PROD=disrupted in schizophrenia 1 /DB_XREF=gi:11037064 /UG=Hs.26985 disrupted in schizophrenia 1 /FL=gb:NM_018662.1 gb:AF222980.1	NM_018662		NP_061132
206099_at	0.003862	gb:NM_006255.1 /DEF=Homo sapiens protein kinase C, eta (PRKCH), mRNA. /FEA=mRNA /GEN=PRKCH /PROD=protein kinase C, eta /DB_XREF=gi:5453971 /UG=Hs.315366 protein kinase C, eta /FL=gb:M55284.1 gb:NM_006255.1	NM_006255		NP_006246

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206111_at	1.26E-04	gb:NM_002934.1 /DEF=Homo sapiens ribonuclease, RNase A family, 2 (liver, eosinophil-derived neurotoxin) (RNASE2), mRNA. /FEA=mRNA /GEN=RNASE2 /PROD=ribonuclease, RNase A family, 2 (liver,eosinophil-derived neurotoxin) /DB_XREF=gi:4506548 /UG=Hs.728 ribonuclease, RNase A family, 2 (liver, eosinophil-derived neurotoxin) /FL=gb:M24157.1 gb:M30510.1 gb:NM_002934.1 gb:M28129.1	NM_002934		NP_002925
206116_s_at	2.75E-04	gb:NM_000366.1 /DEF=Homo sapiens tropomyosin 1 (alpha) (TPM1), mRNA. /FEA=mRNA /GEN=TPM1 /PROD=tropomyosin 1 (alpha) /DB_XREF=gi:4507646 /UG=Hs.77899 tropomyosin 1 (alpha) /FL=gb:M19713.1 gb:NM_000366.1	NM_000366		NP_000357
206132_at	0.0375	gb:NM_002387.1 /DEF=Homo sapiens mutated in colorectal cancers (MCC), mRNA. /FEA=mRNA /GEN=MCC /PROD=mutated in colorectal cancers /DB_XREF=gi:4505128 /UG=Hs.1345 mutated in colorectal cancers /FL=gb:M62397.1 gb:NM_002387.1	NM_002387		NP_002378
206133_at	0.040938	gb:NM_017523.1 /DEF=Homo sapiens XIAP associated factor-1 (HSXIAPAF1), mRNA. /FEA=mRNA /GEN=HSXIAPAF1 /PROD=XIAP associated factor-1 /DB_XREF=gi:8923794 /UG=Hs.139262 XIAP associated factor-1 /FL=gb:NM_017523.1	NM_017523		NP_059993

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206141_at	0.031704	gb:NM_014484.1 /DEF=Homo sapiens molybdopterin synthase sulfurylase (MOCS3), mRNA. /FEA=mRNA /GEN=MOCS3 /PROD=molybdopterin synthase sulfurylase /DB_XREF=gi:7657338 /UG=Hs.159410 molybdopterin synthase sulfurylase /FL=gb:AF102544.1 gb:NM_014484.1	NM_014484		NP_055299
206150_at	5.03E-04	gb:NM_001242.1 /DEF=Homo sapiens tumor necrosis factor receptor superfamily, member 7 (TNFRSF7), mRNA. /FEA=mRNA /GEN=TNFRSF7 /PROD=CD27 antigen /DB_XREF=gi:4507586 /UG=Hs.180841 tumor necrosis factor receptor superfamily, member 7 /FL=gb:M63928.1 gb:NM_001242.1	NM_001242		NP_001233
206157_at	1.36E-04	gb:NM_002852.1 /DEF=Homo sapiens pentaxin-related gene, rapidly induced by IL-1 beta (PTX3), mRNA. /FEA=mRNA /GEN=PTX3 /PROD=pentaxin-related gene, rapidly induced by IL-1beta /DB_XREF=gi:4506332 /UG=Hs.2050 pentaxin-related gene, rapidly induced by IL-1 beta /FL=gb:M31166.1 gb:NM_002852.1	NM_002852		NP_002843
206158_s_at	2.05E-04	gb:NM_003418.1 /DEF=Homo sapiens zinc finger protein 9 (a cellular retroviral nucleic acid binding protein) (ZNF9), mRNA. /FEA=mRNA /GEN=ZNF9 /PROD=zinc finger protein 9 (a cellular retroviral nucleic acid binding protein) /DB_XREF=gi:4827070 /UG=Hs.2110 zinc finger protein 9 (a cellular retroviral nucleic acid binding protein) /FL=gb:NM_003418.1 gb:M28372.1	NM_003418		NP_003409

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206170_at	0.003171	gb:NM_000024.2 /DEF=Homo sapiens adrenergic, beta-2-, receptor, surface (ADRB2), mRNA. /FEA=mRNA /GEN=ADRB2 /PROD=adrenergic, beta-2-, receptor, surface /DB_XREF=gi:13162366 /UG=Hs.2551 adrenergic, beta-2-, receptor, surface /FL=gb:NM_000024.2 gb:M15169.1	NM_000024		NP_000015
206174_s_at	1.26E-04	gb:NM_002721.2 /DEF=Homo sapiens protein phosphatase 6, catalytic subunit (PPP6C), mRNA. /FEA=mRNA /GEN=PPP6C /PROD=protein phosphatase 6, catalytic subunit /DB_XREF=gi:5729987 /UG=Hs.279563 protein phosphatase 6, catalytic subunit /FL=gb:NM_002721.2	NM_002721		NP_002712
206177_s_at	3.26E-04	gb:NM_000045.2 /DEF=Homo sapiens arginase, liver (ARG1), mRNA. /FEA=mRNA /GEN=ARG1 /PROD=arginase, type I /DB_XREF=gi:10947138 /UG=Hs.289057 arginase, liver /FL=gb:NM_000045.2 gb:M14502.1	NM_000045		NP_000036
206183_s_at	0.001116	gb:NM_014606.1 /DEF=Homo sapiens hect domain and RLD 3 (HERC3), mRNA. /FEA=mRNA /GEN=HERC3 /PROD=hect domain and RLD 3 /DB_XREF=gi:7657151 /UG=Hs.35804 hect domain and RLD 3 /FL=gb:D25215.1 gb:NM_014606.1	NM_014606		NP_055421
206200_s_at	3.26E-04	gb:NM_001157.1 /DEF=Homo sapiens annexin A11 (ANXA11), mRNA. /FEA=mRNA /GEN=ANXA11 /PROD=annexin XI /DB_XREF=gi:4557316 /UG=Hs.75510 annexin A11 /FL=gb:NM_001157.1 gb:L19605.1	NM_001157		NP_665876

Gene Identifier	p-value	D scription	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206208_at	0.001108	gb:NM_000717.2 /DEF=Homo sapiens carbonic anhydrase IV (CA4), mRNA. /FEA=mRNA /GEN=CA4 /PROD=carbonic anhydrase IV precursor /DB_XREF=gi:9951925 /UG=Hs.89485 carbonic anhydrase IV /FL=gb:M83670.1 gb:NM_000717.2	NM_000717		NP_000708
206209_s_at	0.041055	gb:NM_000717.2 /DEF=Homo sapiens carbonic anhydrase IV (CA4), mRNA. /FEA=mRNA /GEN=CA4 /PROD=carbonic anhydrase IV precursor /DB_XREF=gi:9951925 /UG=Hs.89485 carbonic anhydrase IV /FL=gb:M83670.1 gb:NM_000717.2	NM_000717		NP_000708
206222_at	0.026891	gb:NM_003841.1 /DEF=Homo sapiens tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain (TNFRSF10C), mRNA. /FEA=mRNA /GEN=TNFRSF10C /PROD=tumor necrosis factor receptor superfamily,member 10c, decoy without an intracellular domain /DB_XREF=gi:10835042 /UG=Hs.119684 tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain /FL=gb:NM_003841.1 gb:AF012629.1 gb:AF020502.1 gb:AF016267.1 gb:AF033854.1 gb:AF014794.1	NM_003841		NP_003832
206235_at	8.68E-04	gb:NM_002312.1 /DEF=Homo sapiens ligase IV, DNA, ATP-dependent (LIG4), mRNA. /FEA=mRNA /GEN=LIG4 /PROD=DNA ligase IV /DB_XREF=gi:4504996 /UG=Hs.166091 ligase IV, DNA, ATP-dependent /FL=gb:NM_002312.1	NM_002312		NP_002303

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206240_s_at	7.58E-04	gb:NM_003437.1 /DEF=Homo sapiens zinc finger protein 136 (clone pHZ-20) (ZNF136), mRNA. /FEA=mRNA /GEN=ZNF136 /PROD=zinc finger protein 136 (clone pHZ-20) /DB_XREF=gi:4507986 /UG=Hs.182828 zinc finger protein 136 (clone pHZ-20) /FL=gb:NM_003437.1	NM_003437		NP_003428
206244_at	0.004798	gb:NM_000573.1 /DEF=Homo sapiens complement component (3b4b) receptor 1, including Knops blood group system (CR1), mRNA. /FEA=mRNA /GEN=CR1 /PROD=complement component (3b4b) receptor 1,including Knops blood group system /DB_XREF=gi:10834973 /UG=Hs.193716 complement component (3b4b) receptor 1, including Knops blood group system /FL=gb:NM_000573.1	NM_000573		NP_000642
206245_s_at	3.26E-04	gb:NM_006469.1 /DEF=Homo sapiens NS1-binding protein (NS1 BP), mRNA. /FEA=mRNA /GEN=NS1-BP /PROD=NS1-binding protein /DB_XREF=gi:5453803 /UG=Hs.197298 NS1-binding protein /FL=gb:NM_006469.1	NM_006469		NP_006460
206267_s_at	0.012681	gb:NM_002378.1 /DEF=Homo sapiens megakaryocyte-associated tyrosine kinase (MATK), mRNA. /FEA=mRNA /GEN=MATK /PROD=megakaryocyte-associated tyrosine kinase /DB_XREF=gi:4505108 /UG=Hs.274 megakaryocyte-associated tyrosine kinase /FL=gb:BC000114.1 gb:BC003109.1 gb:NM_002378.1 gb:L18974.1	NM_002378		NP_647612

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206284_x_at	0.026891	gb:NM_001834.1 /DEF=Homo sapiens clathrin, light polypeptide (Lcb) (CLTB), transcript variant nonbrain, mRNA. /FEA=mRNA /GEN=CLTB /PROD=clathrin, light polypeptide B (Lcb) isoform a /DB_XREF=gi:4502900 /UG=Hs.73919 clathrin, light polypeptide (Lcb) /FL=gb:M20470.1 gb:NM_001834.1	NM_001834		NP_009028
206295_at	0.011725	gb:NM_001562.1 /DEF=Homo sapiens interleukin 18 (interferon-gamma-inducing factor) (IL18), mRNA. /FEA=mRNA /GEN=IL18 /PROD=interleukin 18 /DB_XREF=gi:4504652 /UG=Hs.83077 interleukin 18 (interferon-gamma-inducing factor) /FL=gb:D49950.1 gb:AF077611.1 gb:NM_001562.1	NM_001562		NP_001553
206302_s_at	1.26E-04	gb:NM_019094.1 /DEF=Homo sapiens nudix (nucleoside diphosphate linked moiety X)-type motif 4 (NUDT4), mRNA. /FEA=mRNA /GEN=NUDT4 /PROD=nudix (nucleoside diphosphate linked moietyX)-type motif 4 /DB_XREF=gi:10800135 /UG=Hs.92381 nudix (nucleoside diphosphate linked moiety X)-type motif 4 /FL=gb:NM_019094.1 gb:AF191649.1 gb:AF191650.1 gb:AF191653.1	NM_019094		NP_061967
206332_s_at	0.033381	gb:NM_005531.1 /DEF=Homo sapiens interferon, gamma-inducible protein 16 (IFI16), mRNA. /FEA=mRNA /GEN=IFI16 /PROD=interferon, gamma-inducible protein 16 /DB_XREF=gi:5031778 /UG=Hs.155530 interferon, gamma-inducible protein 16 /FL=gb:M63838.1 gb:NM_005531.1	NM_005531		NP_005522

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206335_at	6.68E-04	gb:NM_000512.2 /DEF=Homo sapiens galactosamine (N-acetyl)-6-sulfate sulfatase (Morquio syndrome, mucopolysaccharidosis type IVA) (GALNS), mRNA. /FEA=mRNA /GEN=GALNS /PROD=N-acetylgalactosamine-6-sulfatase precursor /DB_XREF=gi:9945384 /UG=Hs.159479 galactosamine (N-acetyl)-6-sulfate sulfatase (Morquio syndrome, mucopolysaccharidosis type IVA) /FL=gb:NM_000512.2	NM_000512		NP_000503
206337_at	1.26E-04	gb:NM_001838.1 /DEF=Homo sapiens chemokine (C-C motif) receptor 7 (CCR7), mRNA. /FEA=mRNA /GEN=CCR7 /PROD=chemokine (C-C motif) receptor 7 /DB_XREF=gi:4502640 /UG=Hs.1652 chemokine (C-C motif) receptor 7 /FL=gb:L08176.1 gb:NM_001838.1 gb:L31581.1	NM_001838		NP_001829
206364_at	3.09E-04	gb:NM_014875.1 /DEF=Homo sapiens KIAA0042 gene product (KIAA0042), mRNA. /FEA=mRNA /GEN=KIAA0042 /PROD=KIAA0042 gene product /DB_XREF=gi:7661877 /UG=Hs.3104 KIAA0042 gene product /FL=gb:D26361.1 gb:NM_014875.1	NM_014875		NP_055690
206371_at	0.006096	gb:NM_000804.2 /DEF=Homo sapiens folate receptor 3 (gamma) (FOLR3), mRNA. /FEA=mRNA /GEN=FOLR3 /PROD=folate receptor 3 precursor /DB_XREF=gi:9257219 /UG=Hs.352 folate receptor 3 (gamma) /FL=gb:U08470.1 gb:U08471.1 gb:NM_000804.2	NM_000804		NP_000795

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206380_s_at	1.26E-04	gb:NM_002621.1 /DEF=Homo sapiens properdin P factor, complement (PFC), mRNA. /FEA=mRNA /GEN=PFC /PROD=properdin P factor, complement /DB_XREF=gi:4505736 /UG=Hs.53155 properdin P factor, complement /FL=gb:NM_002621.1 gb:M83652.1	NM_002621		NP_002612
206383_s_at	0.028665	gb:NM_012297.1 /DEF=Homo sapiens Ras-GTPase activating protein SH3 domain-binding protein 2 (KIAA0660), mRNA. /FEA=mRNA /GEN=KIAA0660 /PROD=Ras-GTPase activating protein SH3 domain-bindingprotein 2 /DB_XREF=gi:7110662 /UG=Hs.6727 Ras-GTPase activating protein SH3 domain-binding protein 2 /FL=gb:AF051311.1 gb:AF053535.1 gb:NM_012297.1	NM_012297		NP_036429
206405_x_at	1.26E-04	gb:NM_004505.1 /DEF=Homo sapiens ubiquitin specific protease 6 (Tre-2 oncogene) (USP6), mRNA. /FEA=mRNA /GEN=USP6 /PROD=ubiquitin specific protease 6 /DB_XREF=gi:4758563 /UG=Hs.111065 ubiquitin specific protease 6 (Tre-2 oncogene) /FL=gb:NM_004505.1	NM_004505		NP_004496
206429_at	0.033491	gb:NM_005242.2 /DEF=Homo sapiens coagulation factor II (thrombin) receptor-like 1 (F2RL1), mRNA. /FEA=mRNA /GEN=F2RL1 /PROD=coagulation factor II (thrombin) receptor-like 1precursor /DB_XREF=gi:8051581 /UG=Hs.154299 coagulation factor II (thrombin) receptor-like 1 /FL=gb:U34038.1 gb:NM_005242.2	NM_005242		NP_005233

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206440_at	1.28E-04	gb:NM_004664.1 /DEF=Homo sapiens Vertebrate LIN7 homolog 1, Tax interaction protein 33 (VELI1), mRNA. /FEA=mRNA /GEN=VELI1 /PROD=Vertebrate LIN7 homolog 1, Tax interactionprotein 33 /DB_XREF=gi:4759305 /UG=Hs.178215 Vertebrate LIN7 homolog 1, Tax interaction protein 33 /FL=gb:AF087693.1 gb:NM_004664.1 gb:AF173081.1	NM_004664		NP_004655
206488_s_at	0.016934	gb:NM_000072.1 /DEF=Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin receptor) (CD36), mRNA. /FEA=mRNA /GEN=CD36 /PROD=CD36 antigen (collagen type I receptor,thrombospondin receptor) /DB_XREF=gi:4557418 /UG=Hs.75613 CD36 antigen (collagen type I receptor, thrombospondin receptor) /FL=gb:M24795.1 gb:M98398.1 gb:L06850.1 gb:NM_000072.1	NM_000072		NP_000063
206499_s_at	0.002354	gb:NM_001269.1 /DEF=Homo sapiens chromosome condensation 1 (CHC1), mRNA. /FEA=mRNA /GEN=CHC1 /PROD=chromosome condensation 1 /DB_XREF=gi:4502800 /UG=Hs.84746 chromosome condensation 1 /FL=gb:NM_001269.1	NM_001269		NP_001260
206513_at	0.013223	gb:NM_004833.1 /DEF=Homo sapiens absent in melanoma 2 (AIM2), mRNA. /FEA=mRNA /GEN=AIM2 /PROD=absent in melanoma 2 /DB_XREF=gi:4757733 /UG=Hs.105115 absent in melanoma 2 /FL=gb:AF024714.1 gb:NM_004833.1	NM_004833		NP_004824

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206515_at	0.041055	gb:NM_000896.1 /DEF=Homo sapiens cytochrome P450, subfamily IVF, polypeptide 3 (leukotriene B4 omega hydroxylase) (CYP4F3), mRNA. /FEA=mRNA /GEN=CYP4F3 /PROD=cytochrome P450, subfamily IVF, polypeptide 3 /DB_XREF=gi:4503240 /UG=Hs.106242 cytochrome P450, subfamily IVF, polypeptide 3 (leukotriene B4 omega hydroxylase) /FL=gb:AB002454.1 gb:D12620.1 gb:NM_000896.1	NM_000896		NP_000887
206522_at	2.05E-04	gb:NM_004668.1 /DEF=Homo sapiens maltase-glucoamylase (alpha-glucosidase) (MGAM), mRNA. /FEA=mRNA /GEN=MGAM /PROD=alpha-glucosidase /DB_XREF=gi:4758711 /UG=Hs.122785 maltase-glucoamylase (alpha-glucosidase) /FL=gb:AF016833.1 gb:NM_004668.1	NM_004668		NP_004659
206542_s_at	1.26E-04	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	AV725365	Hs.198296	NP_620614
206544_x_at	0.033381	gb:NM_003070.1 /DEF=Homo sapiens SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2 (SMARCA2), mRNA. /FEA=mRNA /GEN=SMARCA2 /PROD=SWISNF related, matrix associated, actindependent regulator of chromatin, subfamily a, member 2 /DB_XREF=gi:4507068 /UG=Hs.198296 SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2 /FL=gb:NM_003070.1 gb:D26155.1	NM_003070		NP_620614

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206551_x_at	0.004798	gb:NM_017644.1 /DEF=Homo sapiens hypothetical protein FLJ20059 (FLJ20059), mRNA. /FEA=mRNA /GEN=FLJ20059 /PROD=hypothetical protein FLJ20059 /DB_XREF=gi:8923060 /UG=Hs.246875 hypothetical protein FLJ20059 /FL=gb:NM_017644.1	NM_017644		NP_060114
206555_s_at	0.003529	gb:NM_017736.1 /DEF=Homo sapiens hypothetical protein FLJ20274 (FLJ20274), mRNA. /FEA=mRNA /GEN=FLJ20274 /PROD=hypothetical protein FLJ20274 /DB_XREF=gi:8923246 /UG=Hs.268371 hypothetical protein FLJ20274 /FL=gb:BC000448.1 gb:NM_017736.1	NM_017736		NP_060206
206559_x_at	0.010205	gb:NM_001403.1 /DEF=Homo sapiens eukaryotic translation elongation factor 1 alpha 1-like 14 (EEF1A1L14), mRNA. /FEA=mRNA /GEN=EEF1A1L14 /PROD=eukaryotic translation elongation factor 1 alpha1-like 14 /DB_XREF=gi:4503472 /UG=Hs.274466 eukaryotic translation elongation factor 1 alpha 1-like 14 /FL=gb:NM_001403.1	NM_001403		
206567_s_at	0.037567	gb:NM_016436.1 /DEF=Homo sapiens hepatocellular carcinoma-associated antigen 58 (LOC51230), mRNA. /FEA=mRNA /GEN=LOC51230 /PROD=hepatocellular carcinoma-associated antigen 58 /DB_XREF=gi:7705990 /UG=Hs.301055 hepatocellular carcinoma-associated antigen 58 /FL=gb:AF220416.1 gb:NM_016436.1	NM_016436		NP_057520

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206571_s_at	0.001116	gb:NM_004834.1 /DEF=Homo sapiens mitogen-activated protein kinase kinase kinase kinase 4 (MAP4K4), mRNA. /FEA=mRNA /GEN=MAP4K4 /PROD=mitogen-activated protein kinase kinase kinasekinase 4 /DB_XREF=gi:4758523 /UG=Hs.3628 mitogen-activated protein kinase kinase kinase kinase 4 /FL=gb:AF096300.1 gb:NM_004834.1	NM_004834		NP_663720
206576_s_at	0.005852	gb:NM_001712.1 /DEF=Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein) (CEACAM1), mRNA. /FEA=mRNA /GEN=CEACAM1 /PROD=carcinoembryonic antigen-related cell adhesionmolecule 1 (biliary glycoprotein) /DB_XREF=gi:4502404 /UG=Hs.50964 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein) /FL=gb:NM_001712.1	NM_001712		NP_001703
206584_at	0.02145	gb:NM_015364.1 /DEF=Homo sapiens MD-2 protein (MD-2), mRNA. /FEA=mRNA /GEN=MD-2 /PROD=MD-2 protein /DB_XREF=gi:7662503 /UG=Hs.69328 MD-2 protein /FL=gb:AB018549.1 gb:NM_015364.1 gb:AF168121.1	NM_015364		NP_056179
206592_s_at	0.001116	gb:NM_003938.1 /DEF=Homo sapiens adaptor-related protein complex 3, delta 1 subunit (AP3D1), mRNA. /FEA=mRNA /GEN=AP3D1 /PROD=adaptor-related protein complex 3, delta 1subunit /DB_XREF=gi:4501976 /UG=Hs.75056 adaptor-related protein complex 3, delta 1 subunit /FL=gb:U91930.1 gb:NM_003938.1	NM_003938		NP_003929

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206618_at	5.03E-04	gb:NM_003855.1 /DEF=Homo sapiens interleukin 18 receptor 1 (IL18R1), mRNA. /FEA=mRNA /GEN=IL18R1 /PROD=interleukin 18 receptor 1 /DB_XREF=gi:4504654 /UG=Hs.159301 interleukin 18 receptor 1 /FL=gb:U43672.1 gb:NM_003855.1	NM_003855		NP_003846
206649_s_at	0.0174	gb:NM_006521.2 /DEF=Homo sapiens transcription factor binding to IGHM enhancer 3 (TFE3), mRNA. /FEA=mRNA /GEN=TFE3 /PROD=transcription factor binding to IGHM enhancer 3 /DB_XREF=gi:8659573 /UG=Hs.274184 transcription factor binding to IGHM enhancer 3 /FL=gb:NM_006521.2	NM_006521		NP_006512
206652_at	0.011626	gb:NM_016384.1 /DEF=Homo sapiens hypothetical protein (HSPC050), mRNA. /FEA=mRNA /GEN=HSPC050 /PROD=hypothetical protein /DB_XREF=gi:7705438 /UG=Hs.278985 hypothetical protein /FL=gb:AF161535.1 gb:NM_016384.1	NM_016384		
206656_s_at	1.26E-04	gb:BC000353.1 /DEF=Homo sapiens, myogenic factor 3, clone MGC:8630, mRNA, complete cds. /FEA=mRNA /PROD=myogenic factor 3 /DB_XREF=gi:12653170 /UG=Hs.284203 myogenic factor 3 /FL=gb:BC000353.1 gb:NM_002478.2	BC000353		
206672_at	0.004798	gb:NM_000486.2 /DEF=Homo sapiens aquaporin 2 (collecting duct) (AQP2), mRNA. /FEA=mRNA /GEN=AQP2 /PROD=aquaporin 2 /DB_XREF=gi:4755122 /UG=Hs.37025 aquaporin 2 (collecting duct) /FL=gb:NM_000486.2	NM_000486		NP_000477

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206676_at	1.36E-04	gb:M33326.1 /DEF=Human nonspecific cross-reacting antigen (NCA) mRNA, complete cds. /FEA=mRNA /GEN=NCA /PROD=non-specific cross reacting antigen /DB_XREF=gi:189101 /UG=Hs.41 carcinoembryonic antigen-related cell adhesion molecule 8 /FL=gb:M33326.1 gb:NM_001816.1	M33326		NP_001807
206687_s_at	7.58E-04	gb:NM_002831.1 /DEF=Homo sapiens protein tyrosine phosphatase, non-receptor type 6 (PTPN6), mRNA. /FEA=mRNA /GEN=PTPN6 /PROD=protein tyrosine phosphatase, non-receptor type6 /DB_XREF=gi:4506296 /UG=Hs.63489 protein tyrosine phosphatase, non-receptor type 6 /FL=gb:M74903.1 gb:M77273.1 gb:NM_002831.1	NM_002831		NP_536859
206688_s_at	0.003171	gb:NM_006693.1 /DEF=Homo sapiens cleavage and polyadenylation specific factor 4, 30kD subunit (CPSF4), mRNA. /FEA=mRNA /GEN=CPSF4 /PROD=cleavage and polyadenylation specific factor 4,30kD subunit /DB_XREF=gi:5729938 /UG=Hs.6351 cleavage and polyadenylation specific factor 4, 30kD subunit /FL=gb:U79569.1 gb:NM_006693.1	NM_006693		NP_006684
206697_s_at	5.67E-04	gb:NM_005143.1 /DEF=Homo sapiens haptoglobin (HP), mRNA. /FEA=mRNA /GEN=HP /PROD=haptoglobin /DB_XREF=gi:4826761 /UG=Hs.75990 haptoglobin /FL=gb:K00422.1 gb:L29394.1 gb:NM_005143.1	NM_005143		NP_005134

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206698_at	6.77E-04	gb:NM_021083.1 /DEF=Homo sapiens Kell blood group precursor (McLeod phenotype) (XK), mRNA. /FEA=mRNA /GEN=XK /PROD=Kell blood group precursor (McLeod phenotype) /DB_XREF=gi:10835266 /UG=Hs.78919 Kell blood group precursor (McLeod phenotype) /FL=gb:NM_021083.1	NM_021083		NP_066569
206743_s_at	0.021312	gb:NM_001671.1 /DEF=Homo sapiens asialoglycoprotein receptor 1 (ASGR1), mRNA. /FEA=mRNA /GEN=ASGR1 /PROD=asialoglycoprotein receptor 1 /DB_XREF=gi:4502250 /UG=Hs.12056 asialoglycoprotein receptor 1 /FL=gb:M10058.1 gb:NM_001671.1	NM_001671		NP_001662
206756_at	0.004659	gb:NM_019886.1 /DEF=Homo sapiens carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7 (CHST7), mRNA. /FEA=mRNA /GEN=CHST7 /PROD=carbohydrate (N-acetylglucosamine 6-O)sulfotransferase 7 /DB_XREF=gi:9845239 /UG=Hs.138155 carbohydrate (N-acetylglucosamine 6-O)sulfotransferase 7 /FL=gb:AB040711.1 gb:AB037187.1 gb:NM_019886.1	NM_019886		NP_063939
206764_x_at	0.027633	gb:NM_023075.1 /DEF=Homo sapiens hypothetical protein FLJ11585 (FLJ11585), mRNA. /FEA=mRNA /GEN=FLJ11585 /PROD=hypothetical protein FLJ11585 /DB_XREF=gi:12751486 /UG=Hs.154145 hypothetical protein FLJ11585 /FL=gb:NM_023075.1	NM_023075		NP_075563

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206765_at	0.005852	gb:AF153820.1 /DEF=Homo sapiens inwardly-rectifying potassium channel Kir2.1 (KCNJ2) mRNA, complete cds. /FEA=mRNA /GEN=KCNJ2 /PROD=inwardly-rectifying potassium channel Kir2.1 /DB_XREF=gi:8132296 /UG=Hs.1547 potassium inwardly-rectifying channel, subfamily J, member 2 /FL=gb:AF011904.1 gb:AF021139.1 gb:NM_000891.1 gb:U16861.1 gb:U12507.1 gb:U22413.1 gb:AF153820.1 gb:U24055.1	AF153820		NP_000882
206790_s_at	7.58E-04	gb:NM_004545.1 /DEF=Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 1 (7kD, MNLL) (NDUFB1), mRNA. /FEA=mRNA /GEN=NDUFB1 /PROD=NADH dehydrogenase (ubiquinone) 1 betasubcomplex, 1 (7kD, MNLL) /DB_XREF=gi:4758775 /UG=Hs.183435 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 1 (7kD, MNLL) /FL=gb:AF054181.1 gb:NM_004545.1	NM_004545		NP_004536
206792_x_at	0.007779	gb:NM_000923.1 /DEF=Homo sapiens phosphodiesterase 4C, cAMP-specific (dunce (Drosophila)-homolog phosphodiesterase E1) (PDE4C), mRNA. /FEA=mRNA /GEN=PDE4C /PROD=phosphodiesterase 4C, cAMP-specific (dunce(Drosophila)-homolog phosphodiesterase E1) /DB_XREF=gi:4505664 /UG=Hs.189 phosphodiesterase 4C, cAMP-specific (dunce (Drosophila)-homolog phosphodiesterase E1) /FL=gb:NM_000923.1	NM_000923		NP_000914

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206816_s_at	0.001323	gb:NM_012436.1 /DEF=Homo sapiens sperm associated antigen 8 (SPAG8), mRNA. /FEA=mRNA /GEN=SPAG8 /PROD=sperm associated antigen 8 /DB_XREF=gi:6912671 /UG=Hs.256747 sperm associated antigen 8 /FL=gb:U12978.2 gb:NM_012436.1	NM_012436		NP_758516
206828_at	0.013223	gb:NM_003328.1 /DEF=Homo sapiens TXK tyrosine kinase (TXK), mRNA. /FEA=mRNA /GEN=TXK /PROD=TXK tyrosine kinase /DB_XREF=gi:4507742 /UG=Hs.29877 TXK tyrosine kinase /FL=gb:NM_003328.1 gb:L27071.1	NM_003328		NP_003319
206829_x_at	6.68E-04	gb:NM_025189.1 /DEF=Homo sapiens hypothetical protein FLJ13659 (FLJ13659), mRNA. /FEA=mRNA /GEN=FLJ13659 /PROD=hypothetical protein FLJ13659 /DB_XREF=gi:13430887 /UG=Hs.301651 hypothetical protein FLJ13659 /FL=gb:NM_025189.1	NM_025189		NP_079465
206833_s_at	0.030771	gb:NM_001108.1 /DEF=Homo sapiens acylphosphatase 2, muscle type (ACYP2), mRNA. /FEA=mRNA /GEN=ACYP2 /PROD=acylphosphatase 2, muscle type /DB_XREF=gi:4557246 /UG=Hs.331598 acylphosphatase 2, muscle type /FL=gb:NM_001108.1	NM_001108		
206834_at	0.008237	gb:NM_000519.2 /DEF=Homo sapiens hemoglobin, delta (HBD), mRNA. /FEA=mRNA /GEN=HBD /PROD=hemoglobin, delta /DB_XREF=gi:6633803 /UG=Hs.36977 hemoglobin, delta /FL=gb:NM_000519.2	NM_000519		NP_000510

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206845_s_at	0.002704	gb:NM_014771.1 /DEF=Homo sapiens 95 kDa retinoblastoma protein binding protein; KIAA0661 gene product (KIAA0661), mRNA. /FEA=mRNA /GEN=KIAA0661 /PROD=95 kDa retinoblastoma protein binding protein;KIAA0661 gene product /DB_XREF=gi:7662229 /UG=Hs.65238 95 kDa retinoblastoma protein binding protein; KIAA0661 gene product /FL=gb:AB014561.1 gb:NM_014771.1	NM_014771		NP_055586
206848_at	0.003539	gb:NM_006896.1 /DEF=Homo sapiens homeo box A7 (HOXA7), mRNA. /FEA=mRNA /GEN=HOXA7 /PROD=homeo box A7 /DB_XREF=gi:5901971 /UG=Hs.70954 homeo box A7 /FL=gb:AF026397.1 gb:NM_006896.1	NM_006896		
206851_at	0.011725	gb:NM_002935.1 /DEF=Homo sapiens ribonuclease, RNase A family, 3 (eosinophil cationic protein) (RNASE3), mRNA. /FEA=mRNA /GEN=RNASE3 /PROD=ribonuclease, RNase A family, 3 (eosinophilcationic protein) /DB_XREF=gi:4506550 /UG=Hs.73839 ribonuclease, RNase A family, 3 (eosinophil cationic protein) /FL=gb:NM_002935.1 gb:M28128.1	NM_002935		NP_002926
206871_at	0.002536	gb:NM_001972.1 /DEF=Homo sapiens elastase 2, neutrophil (ELA2), mRNA. /FEA=mRNA /GEN=ELA2 /PROD=elastase 2, neutrophil /DB_XREF=gi:4503548 /UG=Hs.99863 elastase 2, neutrophil /FL=gb:M34379.1 gb:NM_001972.1	NM_001972		NP_001963

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206875_s_at	1.26E-04	gb:NM_014720.1 /DEF=Homo sapiens Ste20-related serinethreonine kinase (KIAA0204), mRNA. /FEA=mRNA /GEN=KIAA0204 /PROD=Ste20-related serinethreonine kinase /DB_XREF=gi:7661993 /UG=Hs.105751 Ste20-related serinethreonine kinase /FL=gb:D86959.1 gb:NM_014720.1	NM_014720		NP_055535
206900_x_at	0.0174	gb:NM_021047.1 /DEF=Homo sapiens DNA-binding protein (LOC56242), mRNA. /FEA=mRNA /GEN=LOC56242 /PROD=DNA-binding protein /DB_XREF=gi:10567821 /UG=Hs.137582 DNA-binding protein /FL=gb:NM_021047.1 gb:AF038951.1	NM_021047		NP_066385
206920_s_at	0.002106	gb:NM_001499.1 /DEF=Homo sapiens GLE1 (yeast homolog)-like, RNA export mediator (GLE1L), mRNA. /FEA=mRNA /GEN=GLE1L /PROD=GLE1-like, RNA export mediator /DB_XREF=gi:4557626 /UG=Hs.169363 GLE1 (yeast homolog)-like, RNA export mediator /FL=gb:AF058922.1 gb:NM_001499.1	NM_001499		NP_001490
206934_at	0.00228	gb:NM_006065.1 /DEF=Homo sapiens signal regulatory protein, beta, 1 (SIRP-BETA-1), mRNA. /FEA=mRNA /GEN=SIRP-BETA-1 /PROD=signal regulatory protein, beta, 1 /DB_XREF=gi:5174678 /UG=Hs.194784 signal regulatory protein, beta, 1 /FL=gb:NM_006065.1	NM_006065		NP_006056

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
206937_at	5.64E-04	gb:NM_003126.1 /DEF=Homo sapiens spectrin, alpha, erythrocytic 1 (elliptocytosis 2) (SPTA1), mRNA. /FEA=mRNA /GEN=SPTA1 /PROD=spectrin, alpha, erythrocytic 1 (elliptocytosis2) /DB_XREF=gi:4507188 /UG=Hs.1985 spectrin, alpha, erythrocytic 1 (elliptocytosis 2) /FL=gb:M61877.1 gb:NM_003126.1	NM_003126		NP_003117
206943_at	0.034317	gb:NM_004612.1 /DEF=Homo sapiens transforming growth factor, beta receptor I (activin A receptor type II-like kinase, 53kD) (TGFB1), mRNA. /FEA=mRNA /GEN=TGFB1 /PROD=transforming growth factor, beta receptor I(activin A receptor type II-like kinase, 53kD) /DB_XREF=gi:4759225 /UG=Hs.220 transforming growth factor, beta receptor I (activin A receptor type II-like kinase, 53kD) /FL=gb:L11695.1 gb:NM_004612.1	NM_004612		NP_004603
206956_at	0.030413	gb:NM_000711.1 /DEF=Homo sapiens bone gamma-carboxyglutamate (gla) protein (osteocalcin) (BGLAP), mRNA. /FEA=mRNA /GEN=BGLAP /PROD=bone gamma-carboxyglutamate (gla) protein(osteocalcin) /DB_XREF=gi:4502400 /UG=Hs.2558 bone gamma-carboxyglutamate (gla) protein (osteocalcin) /FL=gb:NM_000711.1	NM_000711		NP_000702

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
206978_at	0.013223	gb:NM_000647.2 /DEF=Homo sapiens chemokine (C-C motif) receptor 2 (CCR2), mRNA. /FEA=mRNA /GEN=CCR2 /PROD=chemokine (C-C motif) receptor 2 /DB_XREF=gi:4827072 /UG=Hs.395 chemokine (C-C motif) receptor 2 /FL=gb:U03882.1 gb:NM_000647.2	NM_000647		NP_000639
206995_x_at	1.28E-04	gb:NM_003693.1 /DEF=Homo sapiens acetyl LDL receptor; SREC=scavenger receptor expressed by endothelial cells (SREC), mRNA. /FEA=mRNA /GEN=SREC /PROD=acetyl LDL receptor; SREC=scavenger receptorexpressed by endothelial cells /DB_XREF=gi:4507202 /UG=Hs.57735 acetyl LDL receptor; SREC /FL=gb:D63483.1 gb:D86864.1 gb:NM_003693.1	NM_003693		NP_003684
207004_at	0.041795	gb:NM_000657.1 /DEF=Homo sapiens B-cell CLLlymphoma 2 (BCL2), nuclear gene encoding mitochondrial protein, transcript variant beta, mRNA. /FEA=mRNA /GEN=BCL2 /PROD=B-cell lymphoma protein 2 beta /DB_XREF=gi:4557356 /UG=Hs.79241 B-cell CLLlymphoma 2 /FL=gb:M13995.1 gb:NM_000657.1	NM_000657		NP_000648
207072_at	0.007779	gb:NM_003853.1 /DEF=Homo sapiens interleukin 18 receptor accessory protein (IL18RAP), mRNA. /FEA=mRNA /GEN=IL18RAP /PROD=interleukin 18 receptor accessory protein /DB_XREF=gi:4504656 /UG=Hs.158315 interleukin 18 receptor accessory protein /FL=gb:AF077346.1 gb:NM_003853.1	NM_003853		NP_003844

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207085_x_at	6.93E-04	gb:NM_006140.1 /DEF=Homo sapiens colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage) (CSF2RA), mRNA. /FEA=mRNA /GEN=CSF2RA /PROD=colony stimulating factor 2 receptor, alpha,low-affinity (granulocyte-macrophage) /DB_XREF=gi:5453626 /UG=Hs.182378 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage) /FL=gb:M73832.1 gb:NM_006140.1	NM_006140		NP_758452
207088_s_at	0.002418	gb:NM_003562.1 /DEF=Homo sapiens solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11 (SLC25A11), mRNA. /FEA=mRNA /GEN=SLC25A11 /PROD=solute carrier family 25 (mitochondrial carrier;oxoglutarate carrier), member 11 /DB_XREF=gi:4506996 /UG=Hs.184877 solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11 /FL=gb:NM_003562.1	NM_003562		NP_003553
207093_s_at	0.00136	gb:NM_002544.1 /DEF=Homo sapiens oligodendrocyte myelin glycoprotein (OMG), mRNA. /FEA=mRNA /GEN=OMG /PROD=oligodendrocyte myelin glycoprotein /DB_XREF=gi:4505502 /UG=Hs.194772 oligodendrocyte myelin glycoprotein /FL=gb:NM_002544.1	NM_002544		NP_002535
207098_s_at	0.010205	gb:NM_017927.1 /DEF=Homo sapiens hypothetical protein FLJ20693 (FLJ20693), mRNA. /FEA=mRNA /GEN=FLJ20693 /PROD=hypothetical protein FLJ20693 /DB_XREF=gi:8923620 /UG=Hs.197877 hypothetical protein FLJ20693 /FL=gb:NM_017927.1	NM_017927		NP_284941

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207111_at	1.26E-04	gb:NM_001974.1 /DEF=Homo sapiens egf-like module containing, mucin-like, hormone receptor-like sequence 1 (EMR1), mRNA. /FEA=mRNA /GEN=EMR1 /PROD=egf-like module containing, mucin-like, hormone receptor-like sequence 1 /DB_XREF=gi:4503564 /UG=Hs.2375 egf-like module containing, mucin-like, hormone receptor-like sequence 1 /FL=gb:NM_001974.1	NM_001974		NP_001965
207117_at	7.10E-04	gb:NM_015852.1 /DEF=Homo sapiens Krueppel-related zinc finger protein (H-plk), mRNA. /FEA=mRNA /GEN=H-plk /PROD=Krueppel-related zinc finger protein /DB_XREF=gi:7705388 /UG=Hs.250693 Krueppel-related zinc finger protein /FL=gb:M55422.1 gb:NM_015852.1	NM_015852		NP_056936
207121_s_at	5.07E-04	gb:NM_002748.1 /DEF=Homo sapiens mitogen-activated protein kinase 6 (MAPK6), mRNA. /FEA=mRNA /GEN=MAPK6 /PROD=mitogen-activated protein kinase 6 /DB_XREF=gi:4506090 /UG=Hs.271980 mitogen-activated protein kinase 6 /FL=gb:L77964.1 gb:NM_002748.1	NM_002748		NP_002739
207122_x_at	0.033381	gb:NM_001054.1 /DEF=Homo sapiens sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2 (SULT1A2), mRNA. /FEA=mRNA /GEN=SULT1A2 /PROD=sulfotransferase family, cytosolic, 1A,phenol-preferring, member 2 /DB_XREF=gi:4507302 /UG=Hs.272462 sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2 /FL=gb:NM_001054.1	NM_001054		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207131_x_at	0.005852	gb:NM_013430.1 /DEF=Homo sapiens gamma-glutamyltransferase 1 (GGT1), transcript variant 3, mRNA. /FEA=mRNA /GEN=GGT1 /PROD=gamma-glutamyltransferase 1 /DB_XREF=gi:9845492 /UG=Hs.284380 gamma-glutamyltransferase 1 /FL=gb:M24903.1 gb:NM_005265.1 gb:NM_013430.1	NM_013430		NP_038347
207152_at	0.041055	gb:NM_006180.1 /DEF=Homo sapiens neurotrophic tyrosine kinase, receptor, type 2 (NTRK2), mRNA. /FEA=mRNA /GEN=NTRK2 /PROD=neurotrophic tyrosine kinase, receptor, type 2 /DB_XREF=gi:5453811 /UG=Hs.47860 neurotrophic tyrosine kinase, receptor, type 2 /FL=gb:U12140.1 gb:NM_006180.1	NM_006180		NP_006171
207157_s_at	1.26E-04	gb:NM_005274.1 /DEF=Homo sapiens guanine nucleotide binding protein (G protein), gamma 5 (GNG5), mRNA. /FEA=mRNA /GEN=GNG5 /PROD=guanine nucleotide binding protein (G protein),gamma 5 /DB_XREF=gi:4885286 /UG=Hs.5322 guanine nucleotide binding protein (G protein), gamma 5 /FL=gb:NM_005274.1	NM_005274		NP_005265
207163_s_at	1.26E-04	gb:NM_005163.1 /DEF=Homo sapiens v-akt murine thymoma viral oncogene homolog 1 (AKT1), mRNA. /FEA=mRNA /GEN=AKT1 /PROD=serinethreonine protein kinase /DB_XREF=gi:4885060 /UG=Hs.71816 v-akt murine thymoma viral oncogene homolog 1 /FL=gb:M63167.1 gb:NM_005163.1	NM_005163		NP_005154

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207168_s_at	5.03E-04	gb:NM_004893.1 /DEF=Homo sapiens H2A histone family, member Y (H2AFY), mRNA. /FEA=mRNA /GEN=H2AFY /PROD=H2A histone family, member Y /DB_XREF=gi:4758495 /UG=Hs.75258 H2A histone family, member Y /FL=gb:AF054174.1 gb:NM_004893.1	NM_004893		NP_613258
207186_s_at	2.05E-04	gb:NM_004459.2 /DEF=Homo sapiens fetal Alzheimer antigen (FALZ), mRNA. /FEA=mRNA /GEN=FALZ /PROD=fetal Alzheimer antigen /DB_XREF=gi:6552329 /UG=Hs.99872 fetal Alzheimer antigen /FL=gb:U05237.1 gb:NM_004459.2	NM_004459		NP_004450
207196_s_at	0.00228	gb:NM_006058.1 /DEF=Homo sapiens Nef-associated factor 1 (NAF1), mRNA. /FEA=mRNA /GEN=NAF1 /PROD=Nef-associated factor 1 /DB_XREF=gi:5174608 /UG=Hs.109281 Nef-associated factor 1 /FL=gb:NM_006058.1	NM_006058		NP_006049
207198_s_at	0.007779	gb:NM_004987.2 /DEF=Homo sapiens LIM and senescent cell antigen-like domains 1 (LIMS1), mRNA. /FEA=mRNA /GEN=LIMS1 /PROD=LIM and senescent cell antigen-like domains 1 /DB_XREF=gi:13518025 /UG=Hs.112378 LIM and senescent cell antigen-like domains 1 /FL=gb:U09284.2 gb:NM_004987.2	NM_004987		NP_004978

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
207205_at	0.001116	gb:NM_001817.1 /DEF=Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 4 (CEACAM4), mRNA. /FEA=mRNA /GEN=CEACAM4 /PROD=carcinoembryonic antigen-related cell adhesionmolecule 4 /DB_XREF=gi:4502796 /UG=Hs.12 carcinoembryonic antigen-related cell adhesion molecule 4 /FL=gb:NM_001817.1	NM_001817		NP_001808
207224_s_at	0.016934	gb:NM_016543.1 /DEF=Homo sapiens D-siglec precursor, (LOC51766), mRNA. /FEA=mRNA /GEN=LOC51766 /PROD=D-siglec precursor, /DB_XREF=gi:7706570 /UG=Hs.149250 D-siglec precursor, /FL=gb:AF178981.1 gb:NM_016543.1	NM_016543		NP_057627
207238_s_at	0.005852	gb:NM_002838.1 /DEF=Homo sapiens protein tyrosine phosphatase, receptor type, C (PTPRC), mRNA. /FEA=mRNA /GEN=PTPRC /PROD=protein tyrosine phosphatase, receptor type, C /DB_XREF=gi:4506306 /UG=Hs.170121 protein tyrosine phosphatase, receptor type, C /FL=gb:NM_002838.1	NM_002838		NP_563580
207266_x_at	2.05E-04	gb:NM_016837.1 /DEF=Homo sapiens RNA binding motif, single stranded interacting protein 1 (RBMS1), transcript variant MSSP-3, mRNA. /FEA=mRNA /GEN=RBMS1 /PROD=RNA binding motif, single stranded interactingprotein 1, isoform e /DB_XREF=gi:8400719 /UG=Hs.241567 RNA binding motif, single stranded interacting protein 1 /FL=gb:NM_016837.1	NM_016837		NP_058523

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207269_at	2.17E-04	gb:NM_001925.1 /DEF=Homo sapiens defensin, alpha 4, corticostatin (DEFA4), mRNA. /FEA=mRNA /GEN=DEFA4 /PROD=defensin, alpha 4, preproprotein /DB_XREF=gi:4503302 /UG=Hs.2582 defensin, alpha 4, corticostatin /FL=gb:NM_001925.1	NM_001925		NP_001916
207275_s_at	0.00228	gb:NM_001995.1 /DEF=Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 1 (FACL1), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=FACL1 /PROD=long-chain fatty-acid-coenzyme A ligase 1 /DB_XREF=gi:4503650 /UG=Hs.278333 fatty-acid-Coenzyme A ligase, long-chain 1 /FL=gb:L09229.1 gb:NM_001995.1	NM_001995		NP_001986
207303_at	0.01185	gb:NM_005020.1 /DEF=Homo sapiens phosphodiesterase 1C, calmodulin-dependent (70kD) (PDE1C), mRNA. /FEA=mRNA /GEN=PDE1C /PROD=phosphodiesterase 1C, calmodulin-dependent(70kD) /DB_XREF=gi:4826893 /UG=Hs.41718 phosphodiesterase 1C, calmodulin-dependent (70kD) /FL=gb:U40371.1 gb:NM_005020.1	NM_005020		NP_005011
207320_x_at	5.03E-04	gb:NM_004602.1 /DEF=Homo sapiens staufer (Drosophila, RNA binding protein) (STAU), transcript variant T4, mRNA. /FEA=mRNA /GEN=STAU /PROD=staufer, isoform a /DB_XREF=gi:4759175 /UG=Hs.6113 staufer (Drosophila, RNA-binding protein) /FL=gb:AF061941.1 gb:NM_004602.1	NM_004602		NP_059348

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207323_s_at	0.003257	gb:NM_002385.1 /DEF=Homo sapiens myelin basic protein (MBP), mRNA. /FEA=mRNA /GEN=MBP /PROD=myelin basic protein /DB_XREF=gi:4505122 /UG=Hs.69547 myelin basic protein /FL=gb:M30516.1 gb:NM_002385.1	NM_002385		NP_002376
207329_at	0.01811	gb:NM_002424.1 /DEF=Homo sapiens matrix metalloproteinase 8 (neutrophil collagenase) (MMP8), mRNA. /FEA=mRNA /GEN=MMP8 /PROD=matrix metalloproteinase 8 preproprotein /DB_XREF=gi:4505220 /UG=Hs.73862 matrix metalloproteinase 8 (neutrophil collagenase) /FL=gb:J05556.1 gb:NM_002424.1	NM_002424		NP_002415
207332_s_at	3.26E-04	gb:NM_003234.1 /DEF=Homo sapiens transferrin receptor (p90, CD71) (TFRC), mRNA. /FEA=mRNA /GEN=TFRC /PROD=transferrin receptor (p90, CD71) /DB_XREF=gi:4507456 /UG=Hs.77356 transferrin receptor (p90, CD71) /FL=gb:NM_003234.1	NM_003234		NP_003225
207335_x_at	0.010205	gb:NM_007100.1 /DEF=Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit e (ATP5I), mRNA. /FEA=mRNA /GEN=ATP5I /PROD=ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit e /DB_XREF=gi:6005716 /UG=Hs.85539 ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit e /FL=gb:D50371.1 gb:NM_007100.1	NM_007100		NP_009031

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207347_at	7.58E-04	gb:NM_000124.1 /DEF=Homo sapiens excision repair cross-complementing rodent repair deficiency, complementation group 6 (ERCC6), mRNA. /FEA=mRNA /GEN=ERCC6 /PROD=excision repair cross-complementing rodent repair deficiency, complementation group 6 /DB_XREF=gi:4557564 /UG=Hs.99924 excision repair cross-complementing rodent repair deficiency, complementation group 6 /FL=gb:L04791.1 gb:NM_000124.1	NM_000124		NP_000115
207350_s_at	0.003171	gb:NM_003762.1 /DEF=Homo sapiens vesicle-associated membrane protein 4 (VAMP4), mRNA. /FEA=mRNA /GEN=VAMP4 /PROD=vesicle-associated membrane protein 4 /DB_XREF=gi:4507862 /UG=Hs.102664 vesicle-associated membrane protein 4 /FL=gb:AF044310.1 gb:NM_003762.1	NM_003762		NP_003753
207358_x_at	0.00228	gb:NM_012090.1 /DEF=Homo sapiens actin binding protein; macrophin (microfilament and actin filament cross-linker protein) (ACF7), mRNA. /FEA=mRNA /GEN=ACF7 /PROD=actin binding protein; macrophin (microfilament and actin filament cross-linker protein) /DB_XREF=gi:10048480 /UG=Hs.108258 actin binding protein; macrophin (microfilament and actin filament cross-linker protein) /FL=gb:NM_012090.1 gb:AF141968.1	NM_012090		NP_149033

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207384_at	0.001109	gb:NM_005091.1 /DEF=Homo sapiens peptidoglycan recognition protein (PGLYRP), mRNA. /FEA=mRNA /GEN=PGLYRP /PROD=peptidoglycan recognition protein /DB_XREF=gi:4827035 /UG=Hs.137583 peptidoglycan recognition protein /FL=gb:AF076483.1 gb:NM_005091.1 gb:AF242517.1	NM_005091		NP_005082
207387_s_at	3.26E-04	gb:NM_000167.1 /DEF=Homo sapiens glycerol kinase (GK), mRNA. /FEA=mRNA /GEN=GK /PROD=glycerol kinase /DB_XREF=gi:4504006 /UG=Hs.1466 glycerol kinase /FL=gb:L13943.1 gb:NM_000167.1	NM_000167		NP_000158
207391_s_at	0.041055	gb:NM_003557.1 /DEF=Homo sapiens phosphatidylinositol-4-phosphate 5-kinase, type I, alpha (PIP5K1A), mRNA. /FEA=mRNA /GEN=PIP5K1A /PROD=phosphatidylinositol-4-phosphate 5-kinase, type I, alpha /DB_XREF=gi:4505814 /UG=Hs.149255 phosphatidylinositol-4-phosphate 5-kinase, type I, alpha /FL=gb:U78575.1 gb:NM_003557.1	NM_003557		NP_003548
207419_s_at	0.04734	gb:NM_002872.2 /DEF=Homo sapiens ras-related C3 botulinum toxin substrate 2 (rho family, small GTP binding protein Rac2) (RAC2), mRNA. /FEA=mRNA /GEN=RAC2 /PROD=ras-related C3 botulinum toxin substrate 2 /DB_XREF=gi:9845512 /UG=Hs.173466 ras-related C3 botulinum toxin substrate 2 (rho family, small GTP binding protein Rac2) /FL=gb:NM_002872.2	NM_002872		NP_002863

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207431_s_at	0.041055	gb:NM_003676.1 /DEF=Homo sapiens degenerative spermatocyte (homolog Drosophila; lipid desaturase) (DEGS), mRNA. /FEA=mRNA /GEN=DEGS /PROD=degenerative spermatocyte /DB_XREF=gi:4505192 /UG=Hs.185973 degenerative spermatocyte (homolog Drosophila; lipid desaturase) /FL=gb:AF002668.1 gb:NM_003676.1	NM_003676		NP_659004
207447_s_at	0.027239	gb:NM_013244.1 /DEF=Homo sapiens UDP-N-acetylglucosamine:a-1,3-D-mannoside beta-1,4-N-acetylglucosaminyltransferase IV-homolog (HGNT-IV-H), mRNA. /FEA=mRNA /GEN=HGNT-IV-H /PROD=UDP-N-acetylglucosamine:a-1,3-D-mannosidebeta-1,4-N-acetylglucosaminyltransferase IV-homolog /DB_XREF=gi:7019406 /UG=Hs.227473 UDP-N-acetylglucosamine:a-1,3-D-mannoside beta-1,4-N-acetylglucosaminyltransferase IV-homolog /FL=gb:AB024729.1 gb:NM_013244.1	NM_013244		NP_037376
207467_x_at	1.26E-04	gb:NM_001750.2 /DEF=Homo sapiens calpastatin (CAST), mRNA. /FEA=mRNA /GEN=CAST /PROD=calpastatin /DB_XREF=gi:5729759 /UG=Hs.279607 calpastatin /FL=gb:U58996.2 gb:NM_001750.2	NM_001750		NP_775085
207490_at	0.02141	gb:NM_025019.1 /DEF=Homo sapiens hypothetical protein FLJ13940 (FLJ13940), mRNA. /FEA=mRNA /GEN=FLJ13940 /PROD=hypothetical protein FLJ13940 /DB_XREF=gi:13376538 /UG=Hs.287610 hypothetical protein FLJ13940 /FL=gb:NM_025019.1	NM_025019		NP_079295

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207498_s_at	0.031424	gb:NM_000106.1 /DEF=Homo sapiens cytochrome P450, subfamily IID (debrisoquine, sparteine, etc., -metabolizing), polypeptide 6 (CYP2D6), mRNA. /FEA=mRNA /GEN=CYP2D6 /PROD=cytochrome P450, subfamily IID (debrisoquine,sparteine, etc., -metabolizing), polypeptide 6 /DB_XREF=gi:4503222 /UG=Hs.301086 cytochrome P450, subfamily IID (debrisoquine, sparteine, etc., -metabolizing), polypeptide 6 /FL=gb:M20403.1 gb:NM_000106.1	NM_000106		NP_000097
207500_at	0.011862	gb:NM_004347.1 /DEF=Homo sapiens caspase 5, apoptosis-related cysteine protease (CASP5), mRNA. /FEA=mRNA /GEN=CASP5 /PROD=caspase 5, apoptosis-related cysteine protease /DB_XREF=gi:4757913 /UG=Hs.3257 caspase 5, apoptosis-related cysteine protease /FL=gb:NM_004347.1 gb:U28015.1	NM_004347		NP_004338
207508_at	0.010205	gb:NM_001689.1 /DEF=Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit c (subunit 9) isoform 3 (ATP5G3), mRNA. /FEA=mRNA /GEN=ATP5G3 /PROD=ATP synthase, H ⁺ transporting, mitochondrial F0complex, subunit c (subunit 9) isoform 3 /DB_XREF=gi:4502300 /UG=Hs.429 ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit c (subunit 9) isoform 3 /FL=gb:U09813.1 gb:NM_001689.1	NM_001689		NP_001680

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207543_s_at	1.26E-04	gb:NM_000917.1 /DEF=Homo sapiens procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide I (P4HA1), mRNA. /FEA=mRNA /GEN=P4HA1 /PROD=procollagen-proline, 2-oxoglutarate4-dioxygenase (proline 4-hydroxylase), alpha polypeptidel /DB_XREF=gi:4505564 /UG=Hs.76768 procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide I /FL=gb:M24486.1 gb:NM_000917.1	NM_000917		NP_000908
207545_s_at	7.58E-04	gb:NM_003744.1 /DEF=Homo sapiens numb (Drosophila) homolog (NUMB), mRNA. /FEA=mRNA /GEN=NUMB /PROD=numb (Drosophila) homolog /DB_XREF=gi:4505478 /UG=Hs.78890 numb (Drosophila) homolog /FL=gb:NM_003744.1 gb:L40393.1	NM_003744		NP_003735
207554_x_at	0.021165	gb:NM_001060.1 /DEF=Homo sapiens thromboxane A2 receptor (TBXA2R), mRNA. /FEA=mRNA /GEN=TBXA2R /PROD=thromboxane A2 receptor /DB_XREF=gi:4507380 /UG=Hs.89887 thromboxane A2 receptor /FL=gb:NM_001060.1 gb:D38081.1 gb:U27325.1	NM_001060		NP_001051
207556_s_at	0.029453	gb:NM_003646.1 /DEF=Homo sapiens diacylglycerol kinase, zeta (104kD) (DGKZ), mRNA. /FEA=mRNA /GEN=DGKZ /PROD=diacylglycerol kinase, zeta (104kD) /DB_XREF=gi:4503316 /UG=Hs.89981 diacylglycerol kinase, zeta (104kD) /FL=gb:U51477.1 gb:NM_003646.1	NM_003646		NP_003637

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207568_at	0.020793	gb:NM_004198.1 /DEF=Homo sapiens cholinergic receptor, nicotinic, alpha polypeptide 6 (CHRNA6), mRNA. /FEA=mRNA /GEN=CHRNA6 /PROD=cholinergic receptor, nicotinic, alphapolypeptide 6 /DB_XREF=gi:4757981 /UG=Hs.103128 cholinergic receptor, nicotinic, alpha polypeptide 6 /FL=gb:U62435.1 gb:NM_004198.1	NM_004198		NP_004189
207573_x_at	0.003171	gb:NM_006476.1 /DEF=Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F1F0, subunit g (ATP5JG), mRNA. /FEA=mRNA /GEN=ATP5JG /PROD=ATP synthase, H ⁺ transporting, mitochondrialF1F0, subunit g /DB_XREF=gi:5453560 /UG=Hs.107476 ATP synthase, H ⁺ transporting, mitochondrial F1F0, subunit g /FL=gb:AF092124.1 gb:NM_006476.1	NM_006476		NP_006467
207574_s_at	1.26E-04	gb:NM_015675.1 /DEF=Homo sapiens growth arrest and DNA-damage-inducible, beta (GADD45B), mRNA. /FEA=mRNA /GEN=GADD45B /PROD=DKFZP566B133 protein /DB_XREF=gi:9945331 /UG=Hs.110571 growth arrest and DNA-damage-inducible, beta /FL=gb:AF090950.1 gb:NM_015675.1	NM_015675		NP_056490
207585_s_at	0.041055	gb:NM_001001.1 /DEF=Homo sapiens ribosomal protein L36a (RPL36A), mRNA. /FEA=mRNA /GEN=RPL36A /PROD=ribosomal protein L36a /DB_XREF=gi:4506650 /UG=Hs.118857 ribosomal protein L36a /FL=gb:M15661.1 gb:NM_001001.1	NM_001001		NP_000992

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207596_at	0.021435	gb:NM_018515.1 /DEF=Homo sapiens hypothetical protein PRO2176 (PRO2176), mRNA. /FEA=mRNA /GEN=PRO2176 /PROD=hypothetical protein PRO2176 /DB_XREF=gi:8924125 /UG=Hs.127648 hypothetical protein PRO2176 /FL=gb:AF119865.1 gb:NM_018515.1	NM_018515		
207598_x_at	0.021448	gb:NM_005431.1 /DEF=Homo sapiens X-ray repair complementing defective repair in Chinese hamster cells 2 (XRCC2), mRNA. /FEA=mRNA /GEN=XRCC2 /PROD=X-ray repair cross complementing protein 2 /DB_XREF=gi:4885656 /UG=Hs.129727 X-ray repair complementing defective repair in Chinese hamster cells 2 /FL=gb:AF035587.1 gb:NM_005431.1	NM_005431		NP_005422
207601_at	0.004341	gb:NM_014465.1 /DEF=Homo sapiens thyroid hormone slfotransferase (ST1B2), mRNA. /FEA=mRNA /GEN=ST1B2 /PROD=thyroid hormone slfotransferase /DB_XREF=gi:7657620 /UG=Hs.129742 thyroid hormone slfotransferase /FL=gb:U95726.1 gb:D89479.1 gb:NM_014465.1	NM_014465		NP_055280
207605_x_at	0.007779	gb:NM_024498.1 /DEF=Homo sapiens zinc finger protein 117 (HPF9) (ZNF117), mRNA. /FEA=mRNA /GEN=ZNF117 /PROD=zinc finger protein 117 (HPF9) /DB_XREF=gi:13374556 /UG=Hs.133011 zinc finger protein 117 (HPF9) /FL=gb:NM_024498.1 gb:M27879.1	NM_024498		NP_078774

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207610_s_at	0.003171	gb:NM_013447.1 /DEF=Homo sapiens egf-like module containing, mucin-like, hormone receptor-like sequence 2 (EMR2), mRNA. /FEA=mRNA /GEN=EMR2 /PROD=egf-like module containing, mucin-like, hormonereceptor-like sequence 2 /DB_XREF=gi:7305024 /UG=Hs.137354 egf-like module containing, mucin-like, hormone receptor-like sequence 2 /FL=gb:AF114491.1 gb:NM_013447.1	NM_013447		NP_690885
207616_s_at	3.26E-04	gb:NM_004180.1 /DEF=Homo sapiens TRAF family member-associated NFKB activator (TANK), mRNA. /FEA=mRNA /GEN=TANK /PROD=TRAF family member-associated NFKB activator /DB_XREF=gi:4759249 /UG=Hs.146847 TRAF family member-associated NFKB activator /FL=gb:U63830.1 gb:NM_004180.1	NM_004180		NP_597841
207618_s_at	0.00228	gb:NM_004328.1 /DEF=Homo sapiens BCS1 (yeast homolog)-like (BCS1L), mRNA. /FEA=mRNA /GEN=BCS1L /PROD=BCS1 (yeast homolog)-like /DB_XREF=gi:4757851 /UG=Hs.150922 BCS1 (yeast homolog)-like /FL=gb:AF026849.1 gb:NM_004328.1	NM_004328		NP_004319
207627_s_at	5.64E-04	gb:NM_005653.1 /DEF=Homo sapiens transcription factor CP2 (TFCP2), mRNA. /FEA=mRNA /GEN=TFCP2 /PROD=transcription factor CP2 /DB_XREF=gi:5032174 /UG=Hs.154970 transcription factor CP2 /FL=gb:U03495.1 gb:NM_005653.1	NM_005653		NP_005644

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207628_s_at	0.02145	gb:NM_017528.1 /DEF=Homo sapiens putative methyltransferase (HASJ4442), mRNA. /FEA=mRNA /GEN=HASJ4442 /PROD=putative methyltransferase /DB_XREF=gi:8923713 /UG=Hs.155020 putative methyltransferase /FL=gb:NM_017528.1	NM_017528		NP_059998
207654_x_at	1.26E-04	gb:NM_001938.1 /DEF=Homo sapiens down-regulator of transcription 1, TBP-binding (negative cofactor 2) (DR1), mRNA. /FEA=mRNA /GEN=DR1 /PROD=down-regulator of transcription 1 /DB_XREF=gi:4503380 /UG=Hs.16697 down-regulator of transcription 1, TBP-binding (negative cofactor 2) /FL=gb:M97388.1 gb:NM_001938.1	NM_001938		NP_001929
207655_s_at	2.05E-04	gb:NM_013314.1 /DEF=Homo sapiens B cell linker protein (SLP65), mRNA. /FEA=mRNA /GEN=SLP65 /PROD=B cell linker protein /DB_XREF=gi:7019534 /UG=Hs.167746 B cell linker protein /FL=gb:AF068180.1 gb:NM_013314.1	NM_013314		NP_037446
207657_x_at	0.010205	gb:NM_002270.1 /DEF=Homo sapiens karyopherin (importin) beta 2 (KPNB2), mRNA. /FEA=mRNA /GEN=KPNB2 /PROD=karyopherin (importin) beta 2 /DB_XREF=gi:4504906 /UG=Hs.168075 karyopherin (importin) beta 2 /FL=gb:U70322.1 gb:NM_002270.1	NM_002270		NP_694858

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207665_at	0.033381	gb:NM_003813.1 /DEF=Homo sapiens a disintegrin and metalloproteinase domain 21 (ADAM21), mRNA. /FEA=mRNA /GEN=ADAM21 /PROD=a disintegrin and metalloproteinase domain 21preproprotein /DB_XREF=gi:11497039 /UG=Hs.178748 a disintegrin and metalloproteinase domain 21 /FL=gb:NM_003813.1	NM_003813		NP_003804
207667_s_at	0.001054	gb:NM_002756.1 /DEF=Homo sapiens mitogen-activated protein kinase kinase 3 (MAP2K3), mRNA. /FEA=mRNA /GEN=MAP2K3 /PROD=mitogen-activated protein kinase kinase 3 /DB_XREF=gi:4506098 /UG=Hs.180533 mitogen-activated protein kinase kinase 3 /FL=gb:NM_002756.1 gb:L36719.1	NM_002756		NP_659732
207674_at	0.02145	gb:NM_002000.1 /DEF=Homo sapiens Fc fragment of IgA, receptor for (FCAR), mRNA. /FEA=mRNA /GEN=FCAR /PROD=Fc fragment of IgA, receptor for /DB_XREF=gi:4503672 /UG=Hs.193122 Fc fragment of IgA, receptor for /FL=gb:D87855.1 gb:D87861.1 gb:NM_002000.1	NM_002000		NP_579814
207677_s_at	0.00228	gb:NM_013416.1 /DEF=Homo sapiens neutrophil cytosolic factor 4 (40kD) (NCF4), transcript variant 2, mRNA. /FEA=mRNA /GEN=NCF4 /PROD=neutrophil cytosolic factor 4 (40kD), isoform 2 /DB_XREF=gi:7382492 /UG=Hs.196352 neutrophil cytosolic factor 4 (40kD) /FL=gb:BC002798.1 gb:AB025219.1 gb:NM_013416.1	NM_013416		NP_038202

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207686_s_at	0.011377	gb:NM_001228.1 /DEF=Homo sapiens caspase 8, apoptosis-related cysteine protease (CASP8), mRNA. /FEA=mRNA /GEN=CASP8 /PROD=caspase 8, apoptosis-related cysteine protease /DB_XREF=gi:4502582 /UG=Hs.19949 caspase 8, apoptosis-related cysteine protease /FL=gb:U60520.1 gb:NM_001228.1	NM_001228		NP_203522
207700_s_at	0.00228	gb:NM_006534.1 /DEF=Homo sapiens nuclear receptor coactivator 3 (NCOA3), mRNA. /FEA=mRNA /GEN=NCOA3 /PROD=nuclear receptor coactivator 3 /DB_XREF=gi:5729725 /UG=Hs.225977 nuclear receptor coactivator 3 /FL=gb:AF036892.1 gb:NM_006534.1	NM_006534		NP_006525
207704_s_at	0.031948	gb:NM_003644.1 /DEF=Homo sapiens growth arrest-specific 7 (GAS7), transcript variant a, mRNA. /FEA=mRNA /GEN=GAS7 /PROD=growth arrest-specific 7 isoform a /DB_XREF=gi:4503920 /UG=Hs.226133 growth arrest-specific 7 /FL=gb:NM_003644.1	NM_003644		NP_005881
207707_s_at	0.002864	gb:NM_030673.1 /DEF=Homo sapiens SEC13 (S. cerevisiae)-like 1 (SEC13L1), mRNA. /FEA=mRNA /GEN=SEC13L1 /PROD=SEC13 (S. cerevisiae)-like 1 /DB_XREF=gi:13491163 /UG=Hs.227949 SEC13 (S. cerevisiae)-like 1 /FL=gb:BC002634.1 gb:NM_030673.1	NM_030673		NP_109598
207719_x_at	0.00161	gb:NM_014812.1 /DEF=Homo sapiens KIAA0470 gene product (KIAA0470), mRNA. /FEA=mRNA /GEN=KIAA0470 /PROD=KARP-1-binding protein /DB_XREF=gi:7662141 /UG=Hs.25132 KIAA0470 gene product /FL=gb:AB022657.1 gb:NM_014812.1	NM_014812		NP_055627

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207728_at	0.003664	gb:NM_018005.1 /DEF=Homo sapiens hypothetical protein FLJ10139 (FLJ10139), mRNA. /FEA=mRNA /GEN=FLJ10139 /PROD=hypothetical protein FLJ10139 /DB_XREF=gi:8922245 /UG=Hs.272210 hypothetical protein FLJ10139 /FL=gb:NM_018005.1	NM_018005		NP_060475
207730_x_at	0.003171	gb:NM_017932.1 /DEF=Homo sapiens hypothetical protein FLJ20700 (FLJ20700), mRNA. /FEA=mRNA /GEN=FLJ20700 /PROD=hypothetical protein FLJ20700 /DB_XREF=gi:8923629 /UG=Hs.272222 hypothetical protein FLJ20700 /FL=gb:NM_017932.1	NM_017932		NP_060402
207782_s_at	0.045608	gb:NM_007319.1 /DEF=Homo sapiens presenilin 1 (Alzheimer disease 3) (PSEN1), transcript variant I-374., mRNA. /FEA=mRNA /GEN=PSEN1 /PROD=presenilin 1 isoform I-374 /DB_XREF=gi:7549814 /UG=Hs.3260 presenilin 1 (Alzheimer disease 3) /FL=gb:U40380.1 gb:NM_007319.1	NM_007319		NP_015558
207783_x_at	0.005852	gb:NM_017627.1 /DEF=Homo sapiens hypothetical protein FLJ20030 (FLJ20030), mRNA. /FEA=mRNA /GEN=FLJ20030 /PROD=hypothetical protein FLJ20030 /DB_XREF=gi:8923031 /UG=Hs.326456 hypothetical protein FLJ20030 /FL=gb:NM_017627.1	NM_017627		
207785_s_at	1.26E-04	gb:NM_015874.1 /DEF=Homo sapiens H-2K binding factor-2 (LOC51580), mRNA. /FEA=mRNA /GEN=LOC51580 /PROD=H-2K binding factor-2 /DB_XREF=gi:7706215 /UG=Hs.327138 H-2K binding factor-2 /FL=gb:D14041.1 gb:NM_015874.1	NM_015874		NP_056958

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207791_s_at	0.026891	gb:NM_004161.1 /DEF=Homo sapiens RAB1, member RAS oncogene family (RAB1), mRNA. /FEA=mRNA /GEN=RAB1 /PROD=RAB1, member RAS oncogene family /DB_XREF=gi:4758987 /UG=Hs.3642 RAB1, member RAS oncogene family /FL=gb:NM_004161.1 gb:M28209.1	NM_004161		NP_004152
207794_at	0.045608	gb:NM_000648.1 /DEF=Homo sapiens chemokine (C-C motif) receptor 2 (CCR2), mRNA. /FEA=mRNA /GEN=CCR2 /PROD=chemokine (C-C motif) receptor 2 /DB_XREF=gi:4757937 /UG=Hs.395 chemokine (C-C motif) receptor 2 /FL=gb:U03905.1 gb:NM_000648.1 gb:D29984.1	NM_000648		NP_000639
207801_s_at	0.00161	gb:NM_014868.1 /DEF=Homo sapiens ring finger protein 10 (RNF10), mRNA. /FEA=mRNA /GEN=RNF10 /PROD=ring finger protein 10 /DB_XREF=gi:7662652 /UG=Hs.5094 ring finger protein 10 /FL=gb:D87451.1 gb:NM_014868.1	NM_014868		NP_055683
207805_s_at	2.05E-04	gb:NM_002813.1 /DEF=Homo sapiens proteasome (prosome, macropain) 26S subunit, non-ATPase, 9 (PSMD9), mRNA. /FEA=mRNA /GEN=PSMD9 /PROD=proteasome (prosome, macropain) 26S subunit, non-ATPase, 9 /DB_XREF=gi:4506234 /UG=Hs.5648 proteasome (prosome, macropain) 26S subunit, non-ATPase, 9 /FL=gb:AB003177.1 gb:NM_002813.1	NM_002813		NP_002804

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207808_s_at	0.001886	gb:NM_000313.1 /DEF=Homo sapiens protein S (alpha) (PROS1), mRNA. /FEA=mRNA /GEN=PROS1 /PROD=protein S (alpha) /DB_XREF=gi:4506116 /UG=Hs.64016 protein S (alpha) /FL=gb:M15036.1 gb:NM_000313.1	NM_000313		NP_000304
207809_s_at	0.007779	gb:NM_001183.1 /DEF=Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump), subunit 1 (ATP6S1), mRNA. /FEA=mRNA /GEN=ATP6S1 /PROD=ATPase, H+ transporting, lysosomal subunit 1 /DB_XREF=gi:4557340 /UG=Hs.6551 ATPase, H+ transporting, lysosomal (vacuolar proton pump), subunit 1 /FL=gb:NM_001183.1	NM_001183		NP_001174
207827_x_at	1.28E-04	gb:L36675.1 /DEF=Human (clone 2-4) synuclein (NACP) mRNA, complete cds. /FEA=mRNA /GEN=NACP /PROD=synuclein /DB_XREF=gi:556213 /UG=Hs.76930 synuclein, alpha (non A4 component of amyloid precursor) /FL=gb:L36675.1 gb:D31839.1 gb:NM_007308.1	L36675		NP_009292
207847_s_at	0.048071	gb:NM_002456.1 /DEF=Homo sapiens mucin 1, transmembrane (MUC1), mRNA. /FEA=mRNA /GEN=MUC1 /PROD=mucin 1, transmembrane /DB_XREF=gi:4505282 /UG=Hs.89603 mucin 1, transmembrane /FL=gb:NM_002456.1	NM_002456		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207857_at	0.001116	gb:NM_006866.1 /DEF=Homo sapiens leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 (LILRA2), mRNA. /FEA=mRNA /GEN=LILRA2 /PROD=leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 /DB_XREF=gi:5803067 /UG=Hs.94498 leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 /FL=gb:U82275.1 gb:NM_006866.1	NM_006866		NP_006857
207922_s_at	0.001116	gb:NM_005882.2 /DEF=Homo sapiens macrophage erythroblast attacher (MAEA), mRNA. /FEA=mRNA /GEN=MAEA /PROD=erythroblast macrophage attacher /DB_XREF=gi:9257203 /UG=Hs.20815 macrophage erythroblast attacher /FL=gb:AF084928.1 gb:NM_005882.2	NM_005882		NP_005873
207943_x_at	0.02145	gb:NM_006718.1 /DEF=Homo sapiens pleiomorphic adenoma gene-like 1 (PLAGL1), transcript variant 2, mRNA. /FEA=mRNA /GEN=PLAGL1 /PROD=pleiomorphic adenoma gene-like 1 isoform 2 /DB_XREF=gi:6031193 /UG=Hs.75825 pleiomorphic adenoma gene-like 1 /FL=gb:NM_006718.1	NM_006718		NP_006709

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207956_x_at	1.26E-04	gb:NM_015928.1 /DEF=Homo sapiens androgen-induced prostate proliferative shutoff associated protein (AS3), mRNA. /FEA=mRNA /GEN=AS3 /PROD=androgen-induced prostate proliferative shutoffassociated protein /DB_XREF=gi:7705287 /UG=Hs.168625 androgen-induced prostate proliferative shutoff associated protein /FL=gb:U95825.2 gb:NM_015928.1	NM_015928		NP_057012
207966_s_at	0.022795	gb:NM_012201.1 /DEF=Homo sapiens Golgi apparatus protein 1 (GLG1), mRNA. /FEA=mRNA /GEN=GLG1 /PROD=Golgi apparatus protein 1 /DB_XREF=gi:6912389 /UG=Hs.78979 Golgi apparatus protein 1 /FL=gb:U28811.1 gb:U64791.1 gb:NM_012201.1	NM_012201		NP_036333
207974_s_at	7.58E-04	gb:NM_006930.1 /DEF=Homo sapiens S-phase kinase-associated protein 1A (p19A) (SKP1A), mRNA. /FEA=mRNA /GEN=SKP1A /PROD=S-phase kinase-associated protein 1A (p19A) /DB_XREF=gi:5902087 /UG=Hs.227950 S-phase kinase-associated protein 1A (p19A) /FL=gb:NM_006930.1 gb:U33760.1	NM_006930		NP_733779
207980_s_at	2.05E-04	gb:NM_006079.1 /DEF=Homo sapiens Cbpp300-interacting transactivator, with GluAsp-rich carboxy-terminal domain, 2 (CITED2), mRNA. /FEA=mRNA /GEN=CITED2 /PROD=Cbpp300-interacting transactivator, withGluAsp-rich carboxy-terminal domain, 2 /DB_XREF=gi:5174416 /UG=Hs.82071 Cbpp300-interacting transactivator, with GluAsp-rich carboxy-terminal domain, 2 /FL=gb:U65093.1 gb:NM_006079.1	NM_006079		NP_006070

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
207986_x_at	0.00228	gb:NM_001915.1 /DEF=Homo sapiens cytochrome b-561 (CYB561), mRNA. /FEA=mRNA /GEN=CYB561 /PROD=cytochrome b-561 /DB_XREF=gi:11993936 /UG=Hs.153028 cytochrome b-561 /FL=gb:NM_001915.1	NM_001915		
207988_s_at	0.005852	gb:NM_005731.1 /DEF=Homo sapiens actin related protein 23 complex, subunit 2 (34 kD) (ARPC2), mRNA. /FEA=mRNA /GEN=ARPC2 /PROD=actin related protein 23 complex, subunit 2 (34kD) /DB_XREF=gi:5031598 /UG=Hs.83583 actin related protein 23 complex, subunit 2 (34 kD) /FL=gb:AF006085.1 gb:NM_005731.1	NM_005731		NP_690601
207992_s_at	1.26E-04	gb:NM_000480.1 /DEF=Homo sapiens adenosine monophosphate deaminase (isoform E) (AMPD3), mRNA. /FEA=mRNA /GEN=AMPD3 /PROD=adenosine monophosphate deaminase (isoform E) /DB_XREF=gi:4502078 /UG=Hs.83918 adenosine monophosphate deaminase (isoform E) /FL=gb:NM_000480.1	NM_000480		NP_000471
207996_s_at	0.02145	gb:NM_004338.1 /DEF=Homo sapiens chromosome 18 open reading frame 1 (C18ORF1), mRNA. /FEA=mRNA /GEN=C18ORF1 /PROD=chromosome 18 open reading frame 1 /DB_XREF=gi:4757883 /UG=Hs.153498 chromosome 18 open reading frame 1 /FL=gb:AF009426.1 gb:NM_004338.1	NM_004338		NP_004329

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208003_s_at	1.26E-04	gb:NM_006599.1 /DEF=Homo sapiens nuclear factor of activated T-cells 5, tonicity-responsive (NFAT5), mRNA. /FEA=mRNA /GEN=NFAT5 /PROD=nuclear factor of activated T-cells 5,tonicity-responsive /DB_XREF=gi:5729944 /UG=Hs.86998 nuclear factor of activated T-cells 5, tonicity-responsive /FL=gb:AB020634.1 gb:AF163836.1 gb:NM_006599.1	NM_006599		
208018_s_at	0.001116	gb:NM_002110.1 /DEF=Homo sapiens hemopoietic cell kinase (HCK), mRNA. /FEA=mRNA /GEN=HCK /PROD=hemopoietic cell kinase /DB_XREF=gi:4504356 /UG=Hs.89555 hemopoietic cell kinase /FL=gb:M16591.1 gb:NM_002110.1	NM_002110		NP_002101
208030_s_at	0.010205	gb:NM_001119.2 /DEF=Homo sapiens adducin 1 (alpha) (ADD1), transcript variant 1, mRNA. /FEA=mRNA /GEN=ADD1 /PROD=adducin 1 (alpha) isoform a /DB_XREF=gi:7710113 /FL=gb:NM_001119.2	NM_001119		
208039_at	7.58E-04	gb:NM_003048.1 /DEF=Homo sapiens solute carrier family 9 (sodiumhydrogen exchanger), isoform 2 (SLC9A2), mRNA. /FEA=mRNA /GEN=SLC9A2 /PROD=solute carrier family 9 (sodiumhydrogenexchanger), isoform 2 /DB_XREF=gi:4507058 /UG=Hs.103132 solute carrier family 9 (sodiumhydrogen exchanger), isoform 2 /FL=gb:NM_003048.1	NM_003048		NP_003039

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208052_x_at	0.034221	gb:NM_001815.1 /DEF=Homo sapiens carcinoembryonic antigen-related cell adhesion molecule 3 (CEACAM3), mRNA. /FEA=mRNA /GEN=CEACAM3 /PROD=carcinoembryonic antigen-related cell adhesionmolecule 3 /DB_XREF=gi:4502792 /UG=Hs.11 carcinoembryonic antigen-related cell adhesion molecule 3 /FL=gb:L00693.1 gb:NM_001815.1	NM_001815		NP_001806
208055_s_at	0.028067	gb:NM_015601.1 /DEF=Homo sapiens DKFZP564G092 protein (DKFZP564G092), mRNA. /FEA=mRNA /GEN=DKFZP564G092 /PROD=DKFZP564G092 protein /DB_XREF=gi:7661611 /UG=Hs.11123 DKFZP564G092 protein /FL=gb:NM_015601.1	NM_015601		NP_056416
208056_s_at	0.00161	gb:NM_005187.2 /DEF=Homo sapiens core-binding factor, runt domain, alpha subunit 2; translocated to, 3 (CBFA2T3), mRNA. /FEA=mRNA /GEN=CBFA2T3 /PROD=core-binding factor, runt domain, alpha subunit2; translocated to, 3 /DB_XREF=gi:7427510 /UG=Hs.110099 core-binding factor, runt domain, alpha subunit 2; translocated to, 3 /FL=gb:NM_005187.2	NM_005187		NP_787127
208066_s_at	7.58E-04	gb:NM_001514.2 /DEF=Homo sapiens general transcription factor IIB (GTF2B), mRNA. /FEA=mRNA /GEN=GTF2B /PROD=general transcription factor IIB /DB_XREF=gi:13435384 /FL=gb:NM_001514.2	NM_001514		NP_001505

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208076_at	0.042536	gb:NM_003539.2 /DEF=Homo sapiens H4 histone family, member B (H4FB), mRNA. /FEA=mRNA /GEN=H4FB /PROD=H4 histone family, member B /DB_XREF=gi:5579463 /UG=Hs.248179 H4 histone family, member B /FL=gb:NM_003539.2	NM_003539		NP_003530
208091_s_at	5.03E-04	gb:NM_030796.1 /DEF=Homo sapiens hypothetical protein DKFZp564K0822 (DKFZP564K0822), mRNA. /FEA=mRNA /GEN=DKFZP564K0822 /PROD=hypothetical protein DKFZp564K0822 /DB_XREF=gi:13540577 /FL=gb:NM_030796.1	NM_030796		NP_110423
208092_s_at	0.016934	gb:NM_030797.1 /DEF=Homo sapiens hypothetical protein DKFZp566A1524 (DKFZP566A1524), mRNA. /FEA=mRNA /GEN=DKFZP566A1524 /PROD=hypothetical protein DKFZp566A1524 /DB_XREF=gi:13540579 /FL=gb:NM_030797.1	NM_030797		NP_110424
208094_s_at	0.048741	gb:NM_030818.1 /DEF=Homo sapiens hypothetical protein MGC10471 (MGC10471), mRNA. /FEA=mRNA /GEN=MGC10471 /PROD=hypothetical protein MGC10471 /DB_XREF=gi:13540613 /FL=gb:NM_030818.1	NM_030818		NP_110445
208103_s_at	1.26E-04	gb:NM_030920.1 /DEF=Homo sapiens hypothetical protein MGC5350 (MGC5350), mRNA. /FEA=mRNA /GEN=MGC5350 /PROD=hypothetical protein MGC5350 /DB_XREF=gi:13569878 /FL=gb:NM_030920.1	NM_030920		NP_112182

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208114_s_at	0.026891	gb:NM_030980.1 /DEF=Homo sapiens hypothetical protein FLJ12671 (FLJ12671), mRNA. /FEA=mRNA /GEN=FLJ12671 /PROD=hypothetical protein FLJ12671 /DB_XREF=gi:13569959 /FL=gb:NM_030980.1	NM_030980		NP_112242
208120_x_at	5.03E-04	gb:NM_031221.1 /DEF=Homo sapiens hypothetical protein FKSG63 (FKSG63), mRNA. /FEA=mRNA /GEN=FKSG63 /PROD=hypothetical protein FKSG63 /DB_XREF=gi:13654297 /FL=gb:NM_031221.1	NM_031221		
208130_s_at	0.029453	gb:NM_030984.1 /DEF=Homo sapiens thromboxane A synthase 1 (platelet, cytochrome P450, subfamily V) (TBXAS1), transcript variant TXS-II, mRNA. /FEA=mRNA /GEN=TBXAS1 /PROD=thromboxane A synthase 1 (platelet, cytochrome P450, subfamily V), isoform TXS-II /DB_XREF=gi:13699839 /FL=gb:NM_030984.1	NM_030984		NP_112246
208137_x_at	1.26E-04	gb:NM_030972.1 /DEF=Homo sapiens hypothetical protein MGC5384 (MGC5384), mRNA. /FEA=mRNA /GEN=MGC5384 /PROD=hypothetical protein MGC5384 /DB_XREF=gi:13775165 /FL=gb:NM_030972.1	NM_030972		NP_112234
208152_s_at	0.00228	gb:NM_004728.1 /DEF=Homo sapiens DEADH (Asp-Glu-Ala-AspHis) box polypeptide 21 (DDX21), mRNA. /FEA=mRNA /GEN=DDX21 /PROD=DEADH (Asp-Glu-Ala-AspHis) box polypeptide 21 /DB_XREF=gi:13787208 /FL=gb:NM_004728.1	NM_004728		NP_004719

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208184_s_at	0.026891	gb:NM_003274.1 /DEF=Homo sapiens transmembrane protein 1 (TMEM1), mRNA. /FEA=mRNA /GEN=TMEM1 /PROD=transmembrane protein 1 /DB_XREF=gi:4507550 /UG=Hs.94479 transmembrane protein 1 /FL=gb:NM_003274.1 gb:U19252.1	NM_003274		NP_003265
208195_at	0.008036	gb:NM_003319.1 /DEF=Homo sapiens titin (TTN), mRNA. /FEA=mRNA /GEN=TTN /PROD=titin /DB_XREF=gi:4507720 /UG=Hs.172004 titin /FL=gb:NM_003319.1	NM_003319		
208200_at	0.00228	gb:NM_000575.1 /DEF=Homo sapiens interleukin 1, alpha (IL1A), mRNA. /FEA=mRNA /GEN=IL1A /PROD=interleukin 1, alpha /DB_XREF=gi:13236493 /UG=Hs.1722 interleukin 1, alpha /FL=gb:NM_000575.1 gb:M28983.1	NM_000575		NP_000566
208238_x_at	0.001303	gb:NM_013344.1 /DEF=Homo sapiens leucine zipper-like protein (LZLP), mRNA. /FEA=mRNA /GEN=LZLP /PROD=leucine zipper-like protein /DB_XREF=gi:7106350 /UG=Hs.278952 leucine zipper-like protein /FL=gb:AF159055.1 gb:NM_013344.1	NM_013344		NP_037476
208246_x_at	0.003171	gb:NM_017618.1 /DEF=Homo sapiens hypothetical protein FLJ20006 (FLJ20006), mRNA. /FEA=mRNA /GEN=FLJ20006 /PROD=hypothetical protein FLJ20006 /DB_XREF=gi:8923014 /UG=Hs.279007 hypothetical protein FLJ20006 /FL=gb:NM_017618.1	NM_017618		NP_060088

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208248_x_at	3.26E-04	gb:NM_001642.1 /DEF=Homo sapiens amyloid beta (A4) precursor-like protein 2 (APLP2), mRNA. /FEA=mRNA /GEN=APLP2 /PROD=amyloid beta (A4) precursor-like protein 2 /DB_XREF=gi:4502146 /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2 /FL=gb:L09209.1 gb:NM_001642.1 gb:L27631.1 gb:AF168956.1	NM_001642		NP_001633
208262_x_at	0.035773	gb:NM_000243.1 /DEF=Homo sapiens Mediterranean fever (MEFV), mRNA. /FEA=mRNA /GEN=MEFV /PROD=Mediterranean fever protein /DB_XREF=gi:4557742 /UG=Hs.173730 Mediterranean fever /FL=gb:AF018080.1 gb:NM_000243.1	NM_000243		NP_000234
208270_s_at	0.004341	gb:NM_020216.2 /DEF=Homo sapiens arginyl aminopeptidase (aminopeptidase B) (RNPEP), mRNA. /FEA=mRNA /GEN=RNPEP /PROD=arginyl aminopeptidase (aminopeptidase B) /DB_XREF=gi:13443030 /UG=Hs.283667 arginyl aminopeptidase (aminopeptidase B) /FL=gb:NM_020216.2	NM_020216		NP_064601
208281_x_at	0.041795	gb:NM_020364.1 /DEF=Homo sapiens deleted in azoospermia 3 (DAZ3), mRNA. /FEA=mRNA /GEN=DAZ3 /PROD=deleted in azoospermia 3 /DB_XREF=gi:11036657 /UG=Hs.283812 deleted in azoospermia 3 /FL=gb:NM_020364.1 gb:AF248481.1	NM_020364		NP_065097

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208284_x_at	5.03E-04	gb:NM_013421.1 /DEF=Homo sapiens gamma-glutamyltransferase 1 (GGT1), transcript variant 2, mRNA. /FEA=mRNA /GEN=GGT1 /PROD=gamma-glutamyltransferase 1 precursor /DB_XREF=gi:9845490 /UG=Hs.284380 gamma-glutamyltransferase 1 /FL=gb:J04131.1 gb:NM_013421.1	NM_013421		NP_038347
208290_s_at	1.26E-04	gb:NM_001969.1 /DEF=Homo sapiens eukaryotic translation initiation factor 5 (EIF5), mRNA. /FEA=mRNA /GEN=EIF5 /PROD=eukaryotic translation initiation factor 5 /DB_XREF=gi:4503542 /UG=Hs.286236 eukaryotic translation initiation factor 5 /FL=gb:U49436.1 gb:NM_001969.1	NM_001969		NP_001960
208296_x_at	0.004341	gb:NM_014350.1 /DEF=Homo sapiens TNF-induced protein (GG2-1), mRNA. /FEA=mRNA /GEN=GG2-1 /PROD=TNF-induced protein /DB_XREF=gi:7657123 /UG=Hs.17839 TNF-induced protein /FL=gb:AF099936.1 gb:NM_014350.1	NM_014350		NP_055165
208306_x_at	0.041055	gb:NM_021983.2 /DEF=Homo sapiens major histocompatibility complex, class II, DR beta 4 (HLA-DRB4), mRNA. /FEA=mRNA /GEN=HLA-DRB4 /PROD=major histocompatibility complex, class II, DRbeta 4 /DB_XREF=gi:11875206 /UG=Hs.293934 major histocompatibility complex, class II, DR beta 4 /FL=gb:NM_021983.2	NM_021983		NP_068818

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208308_s_at	0.007779	gb:NM_000175.1 /DEF=Homo sapiens glucose phosphate isomerase (GPI), mRNA. /FEA=mRNA /GEN=GPI /PROD=glucose phosphate isomerase /DB_XREF=gi:4504086 /UG=Hs.180532 glucose phosphate isomerase /FL=gb:BC004982.1 gb:K03515.1 gb:NM_000175.1	NM_000175		NP_000166
208310_s_at	5.03E-04	gb:NM_007085.1 /DEF=Homo sapiens follistatin-like 1 (FSTL1), mRNA. /FEA=mRNA /GEN=FSTL1 /PROD=follistatin-like 1 /DB_XREF=gi:5901955 /UG=Hs.296267 follistatin-like 1 /FL=gb:D89937.1 gb:NM_007085.1	NM_007085		NP_009016
208315_x_at	0.001205	gb:NM_003300.1 /DEF=Homo sapiens TNF receptor-associated factor 3 (TRAF3), mRNA. /FEA=mRNA /GEN=TRAF3 /PROD=TNF receptor-associated factor 3 /DB_XREF=gi:4507678 /UG=Hs.297660 TNF receptor-associated factor 3 /FL=gb:NM_003300.1 gb:U21092.1	NM_003300		NP_663778
208319_s_at	0.001116	gb:NM_006743.1 /DEF=Homo sapiens RNA binding motif protein 3 (RBM3), mRNA. /FEA=mRNA /GEN=RBM3 /PROD=RNA binding motif protein 3 /DB_XREF=gi:5803136 /UG=Hs.301404 RNA binding motif protein 3 /FL=gb:NM_006743.1 gb:U28686.1	NM_006743		NP_006734
208325_s_at	0.008036	gb:NM_006738.1 /DEF=Homo sapiens lymphoid blast crisis oncogene (LBC), mRNA. /FEA=mRNA /GEN=LBC /PROD=lymphoid blast crisis oncogene /DB_XREF=gi:5803057 /UG=Hs.301946 lymphoid blast crisis oncogene /FL=gb:U03634.1 gb:NM_006738.1	NM_006738		NP_658913

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208353_x_at	0.002536	gb:NM_020480.1 /DEF=Homo sapiens ankyrin 1, erythrocytic (ANK1), transcript variant 7, mRNA. /FEA=mRNA /GEN=ANK1 /PROD=ankyrin 1, isoform 7 /DB_XREF=gi:10947047 /UG=Hs.183805 ankyrin 1, erythrocytic /FL=gb:NM_020480.1	NM_020480		NP_065214
208370_s_at	0.003862	gb:NM_004414.2 /DEF=Homo sapiens Down syndrome critical region gene 1 (DSCR1), mRNA. /FEA=mRNA /GEN=DSCR1 /PROD=Down syndrome critical region protein 1 /DB_XREF=gi:7657042 /UG=Hs.184222 Down syndrome critical region gene 1 /FL=gb:U28833.2 gb:NM_004414.2	NM_004414		NP_004405
208405_s_at	0.00228	gb:NM_006016.1 /DEF=Homo sapiens CD164 antigen, sialomucin (CD164), mRNA. /FEA=mRNA /GEN=CD164 /PROD=CD164 antigen, sialomucin /DB_XREF=gi:5174406 /UG=Hs.43910 CD164 antigen, sialomucin /FL=gb:D14043.1 gb:NM_006016.1	NM_006016		NP_006007
208420_x_at	0.022795	gb:NM_003170.1 /DEF=Homo sapiens suppressor of Ty (S.cerevisiae) 6 homolog (SUPT6H), mRNA. /FEA=mRNA /GEN=SUPT6H /PROD=suppressor of Ty (S.cerevisiae) 6 homolog /DB_XREF=gi:11321572 /UG=Hs.12303 suppressor of Ty (S.cerevisiae) 6 homolog /FL=gb:NM_003170.1 gb:U46691.1	NM_003170		NP_003161

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208438_s_at	7.58E-04	gb:NM_005248.1 /DEF=Homo sapiens Gardner-Rasheed feline sarcoma viral (v-fgr) oncogene homolog (FGR), mRNA. /FEA=mRNA /GEN=FGR /PROD=Gardner-Rasheed feline sarcoma viral (v-fgr)oncogene homolog /DB_XREF=gi:4885234 /UG=Hs.1422 Gardner-Rasheed feline sarcoma viral (v-fgr) oncogene homolog /FL=gb:M19722.1 gb:NM_005248.1	NM_005248		NP_005239
208442_s_at	0.006705	gb:NM_000051.1 /DEF=Homo sapiens ataxia telangiectasia mutated (includes complementation groups A, C and D) (ATM), mRNA. /FEA=mRNA /GEN=ATM /PROD=ataxia telangiectasia mutated (includes complementation groups A, C and D) /DB_XREF=gi:4502266 /UG=Hs.194382 ataxia telangiectasia mutated (includes complementation groups A, C and D) /FL=gb:U33841.1 gb:NM_000051.1	NM_000051		NP_612150
208454_s_at	0.026891	gb:NM_016134.1 /DEF=Homo sapiens aminopeptidase (LOC51670), mRNA. /FEA=CDS /GEN=LOC51670 /PROD=aminopeptidase /DB_XREF=gi:7706386 /UG=Hs.278993 aminopeptidase /FL=gb:AF107834.1 gb:NM_016134.1	NM_016134		NP_057218
208470_s_at	0.034317	gb:NM_020995.1 /DEF=Homo sapiens haptoglobin-related protein (HPR), mRNA. /FEA=CDS /GEN=HPR /PROD=haptoglobin-related protein /DB_XREF=gi:10337588 /UG=Hs.328822 haptoglobin-related protein /FL=gb:NM_020995.1	NM_020995		NP_066275

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208490_x_at	7.31E-04	gb:NM_003522.1 /DEF=Homo sapiens H2B histone family, member G (H2BFG), mRNA. /FEA=CDS /GEN=H2BFG /PROD=H2B histone family, member G /DB_XREF=gi:4504264 /UG=Hs.182137 H2B histone family, member G /FL=gb:NM_003522.1	NM_003522		NP_003513
208503_s_at	0.027792	gb:NM_021167.1 /DEF=Homo sapiens hypothetical protein WUGSC:H_RG083M05.2 (LOC57798), mRNA. /FEA=CDS /GEN=LOC57798 /PROD=hypothetical protein WUGSC:H_RG083M05.2 /DB_XREF=gi:10863962 /UG=Hs.296445 hypothetical protein WUGSC:H_RG083M05.2 /FL=gb:NM_021167.1	NM_021167		NP_066990
208527_x_at	7.58E-04	gb:NM_003523.1 /DEF=Homo sapiens H2B histone family, member H (H2BFH), mRNA. /FEA=CDS /GEN=H2BFH /PROD=H2B histone family, member H /DB_XREF=gi:4504266 /UG=Hs.182138 H2B histone family, member H /FL=gb:NM_003523.1	NM_003523		NP_003514
208540_x_at	5.03E-04	gb:NM_021039.1 /DEF=Homo sapiens S100 calcium-binding protein A14 (calgizzarin) (S100A14), mRNA. /FEA=CDS /GEN=S100A14 /PROD=S100 calcium-binding protein A14 (calgizzarin) /DB_XREF=gi:10567825 /UG=Hs.247697 S100 calcium-binding protein A14 (calgizzarin) /FL=gb:NM_021039.1	NM_021039		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208546_x_at	9.79E-04	gb:NM_003524.1 /DEF=Homo sapiens H2B histone family, member J (H2BFJ), mRNA. /FEA=CDS /GEN=H2BFJ /PROD=H2B histone family, member J /DB_XREF=gi:4504268 /UG=Hs.249216 H2B histone family, member J /FL=gb:NM_003524.1	NM_003524		NP_003515
208579_x_at	2.05E-04	gb:NM_017445.1 /DEF=Homo sapiens H2B histone family, member S (H2BFS), mRNA. /FEA=CDS /GEN=H2BFS /PROD=H2B histone family, member S /DB_XREF=gi:11036645 /UG=Hs.306200 H2B histone family, member S /FL=gb:NM_017445.1	NM_017445		
208610_s_at	1.26E-04	serine/arginine repetitive matrix 2	AI655799	Hs.197114	NP_057417
208611_s_at	0.02145	gb:U83867.1 /DEF=Human alpha II spectrin mRNA, complete cds. /FEA=mRNA /PROD=alpha II spectrin /DB_XREF=gi:1805279 /UG=Hs.77196 spectrin, alpha, non-erythrocytic 1 (alpha-fodrin) /FL=gb:J05243.1 gb:U83867.1 gb:NM_003127.1	U83867		NP_003118
208615_s_at	5.03E-04	protein tyrosine phosphatase type IVA, member 2	BF795101	Hs.82911	NP_536317
208616_s_at	0.013223	gb:U48297.1 /DEF=Homo sapiens protein tyrosine phosphatase PTPCAAX2 (hPTPCAAX2) mRNA, complete cds. /FEA=mRNA /GEN=hPTPCAAX2 /PROD=protein tyrosine phosphatase PTPCAAX2 /DB_XREF=gi:1777756 /UG=Hs.82911 protein tyrosine phosphatase type IVA, member 2 /FL=gb:U48297.1 gb:NM_003479.1 gb:AF208850.1	U48297		NP_536317

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208620_at	1.26E-04	gb:U24223.1 /DEF=Human alpha-CP1 mRNA, complete cds. /FEA=mRNA /GEN=alpha-CP1 /PROD=alpha-CP1 /DB_XREF=gi:1215670 /UG=Hs.2853 poly(rC)-binding protein 1 /FL=gb:U24223.1 gb:NM_006196.1	U24223		NP_006187
208628_s_at	0.007779	gb:BC002411.1 /DEF=Homo sapiens, Similar to Y box protein 1, clone MGC:8655, mRNA, complete cds. /FEA=mRNA /PROD=Similar to Y box protein 1 /DB_XREF=gi:12803206 /UG=Hs.74497 nuclease sensitive element binding protein 1 /FL=gb:BC002411.1	BC002411		NP_004550
208630_at	0.022758	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit	BG472176	Hs.75860	NP_000173
208631_s_at	0.041055	gb:U04627.1 /DEF=Human 78 kDa gastrin-binding protein mRNA, complete cds. /FEA=mRNA /PROD=78 kDa gastrin-binding protein /DB_XREF=gi:595266 /UG=Hs.75860 hydroxyacyl-Coenzyme A dehydrogenase3-ketoacyl-Coenzyme A thiolaseenoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit /FL=gb:NM_000182.1 gb:D16480.1 gb:U04627.1	U04627		NP_000173
208632_at	1.26E-04	ring finger protein 10	AL578551	Hs.5094	NP_055683
208636_at	3.26E-04	actinin, alpha 1	AI082078	Hs.119000	NP_001093
208640_at	3.26E-04	ras-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1)	BG292367	Hs.173737	NP_061485

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208641_s_at	3.26E-04	gb:BC004247.1 /DEF=Homo sapiens, ras-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1), clone MGC:10547, mRNA, complete cds. /FEA=mRNA /PROD=ras-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1) /DB_XREF=gi:13279010 /UG=Hs.173737 ras-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1) /FL=gb:BC004247.1 gb:M29870.1 gb:M31467.1 gb:NM_006908.2	BC004247		NP_061485
208642_s_at	1.26E-04	X-ray repair complementing defective repair in Chinese hamster cells 5 (double-strand-break rejoining; Ku autoantigen, 80kDa)	AA205834		NP_066964
208643_s_at	1.26E-04	gb:J04977.1 /DEF=Human Ku autoimmune antigen gene, complete cds. /FEA=mRNA /GEN=G22P1 /DB_XREF=gi:186791 /UG=Hs.84981 X-ray repair complementing defective repair in Chinese hamster cells 5 (double-strand-break rejoining; Ku autoantigen, 80kD) /FL=gb:NM_021141.2 gb:J04977.1 gb:M30938.1	J04977	Hs.84981	NP_066964
208645_s_at	5.03E-04	gb:AF116710.1 /DEF=Homo sapiens PRO2640 mRNA, complete cds. /FEA=mRNA /PROD=PRO2640 /DB_XREF=gi:7959918 /UG=Hs.244621 ribosomal protein S14 /FL=gb:BC001126.1 gb:BC003401.1 gb:NM_005617.1 gb:AF116710.1	AF116710		NP_005608

Gen Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208646_at	0.026891	gb:AF116710.1 /DEF=Homo sapiens PRO2640 mRNA, complete cds. /FEA=mRNA /PROD=PRO2640 /DB_XREF=gi:7959918 /UG=Hs.244621 ribosomal protein S14 /FL=gb:BC001126.1 gb:BC003401.1 gb:NM_005617.1 gb:AF116710.1	AF116710		NP_005608
208647_at	7.58E-04	farnesyl-diphosphate farnesyltransferase 1	AA872727	Hs.48876	NP_004453
208648_at	0.02145	valosin-containing protein	W60953	Hs.106357	NP_009057
208650_s_at	0.003368	CD24 antigen (small cell lung carcinoma cluster 4 antigen)	BG327863	Hs.381003	NP_037362
208651_x_at	1.26E-04	gb:M58664.1 /DEF=Homo sapiens CD24 signal transducer mRNA, complete cds. /FEA=mRNA /PROD=signal transducer CD24 /DB_XREF=gi:180167 /UG=Hs.286124 CD24 antigen (small cell lung carcinoma cluster 4 antigen) /FL=gb:M58664.1 gb:L33930.1 gb:NM_013230.1	M58664		NP_037362
208652_at	1.26E-04	gb:BC000400.1 /DEF=Homo sapiens, protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform, clone MGC:8533, mRNA, complete cds. /FEA=mRNA /PROD=protein phosphatase 2 (formerly 2A), catalyticsubunit, alpha isoform /DB_XREF=gi:12653262 /UG=Hs.91773 protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform /FL=gb:BC000400.1 gb:BC002657.1 gb:J03804.1 gb:M36951.1 gb:NM_002715.1	BC000400		NP_002706
208653_s_at	0.033381	Consensus includes gb:AF263279.1 /DEF=Homo sapiens CD164 mRNA, complete cds. /FEA=CDS /PROD=CD164 /DB_XREF=gi:9230740 /UG=Hs.43910 CD164 antigen, sialomucin /FL=gb:AF299341.1 gb:AF299343.1 gb:AF263279.1	AF299343		NP_006007

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208654_s_at	1.26E-04	Consensus includes gb:BF669455 /FEA=EST /DB_XREF=gi:11943350 /DB_XREF=est:602120159F1 /CLONE=IMAGE:4277366 /UG=Hs.43910 CD164 antigen, sialomucin /FL=gb:AF299341.1 gb:AF299343.1 gb:AF263279.1	AF299343		NP_006007
208656_s_at	0.033381	gb:AF135162.1 /DEF=Homo sapiens cyclin I (CYC1) mRNA, complete cds. /FEA=mRNA /GEN=CYC1 /PROD=cyclin I /DB_XREF=gi:7259481 /UG=Hs.79933 cyclin I /FL=gb:D50310.1 gb:BC000420.1 gb:BC004975.1 gb:NM_006835.1 gb:AF135162.1	AF135162		NP_006826
208658_at	0.007779	gb:BC000425.1 /DEF=Homo sapiens, protein disulfide isomerase related protein (calcium-binding protein, intestinal-related), clone MGC:8346, mRNA, complete cds. /FEA=mRNA /PROD=protein disulfide isomerase related protein(calcium-binding protein, intestinal-related) /DB_XREF=gi:12653312 /UG=Hs.93659 protein disulfide isomerase related protein (calcium-binding protein, intestinal-related) /FL=gb:BC000425.1 gb:BC001928.1	BC000425		NP_004902
208659_at	0.005852	gb:AF034607.1 /DEF=Homo sapiens chloride channel ABP mRNA, complete cds. /FEA=mRNA /PROD=chloride channel ABP /DB_XREF=gi:4426566 /UG=Hs.74276 chloride intracellular channel 1 /FL=gb:U93205.1 gb:AF034607.1 gb:AF109197.1 gb:NM_001288.2	AF034607		NP_001279

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208660_at	7.58E-04	gb:BC000105.1 /DEF=Homo sapiens, Similar to CG14740 gene product, clone MGC:2503, mRNA, complete cds. /FEA=mRNA /PROD=Similar to CG14740 gene product /DB_XREF=gi:12652712 /UG=Hs.239760 citrate synthase /FL=gb:BC000105.1 gb:AF047042.1 gb:NM_004077.1	BC000105		NP_004068
208661_s_at	0.018693	Consensus includes gb:AW510696 /FEA=EST /DB_XREF=gi:7148774 /DB_XREF=est:hc89f12.x1 /CLONE=IMAGE:2907215 /UG=Hs.118174 tetratricopeptide repeat domain 3 /FL=gb:D84294.1	D84294		NP_003307
208662_s_at	1.26E-04	Consensus includes gb:AI885338 /FEA=EST /DB_XREF=gi:5590502 /DB_XREF=est:wI92e09.x1 /CLONE=IMAGE:2432392 /UG=Hs.118174 tetratricopeptide repeat domain 3 /FL=gb:D84294.1	D84294		NP_003307
208663_s_at	0.027792	Consensus includes gb:AI652848 /FEA=EST /DB_XREF=gi:4736827 /DB_XREF=est:wb40a04.x1 /CLONE=IMAGE:2308110 /UG=Hs.118174 tetratricopeptide repeat domain 3 /FL=gb:D84294.1	D84294		NP_003307
208668_x_at	1.26E-04	gb:BC003689.1 /DEF=Homo sapiens, high-mobility group (nonhistone chromosomal) protein 17, clone MGC:5301, mRNA, complete cds. /FEA=mRNA /PROD=high-mobility group (nonhistone chromosomal)protein 17 /DB_XREF=gi:13277559 /UG=Hs.181163 high-mobility group (nonhistone chromosomal) protein 17 /FL=gb:BC003689.1 gb:M12623.1	BC003689		NP_005508

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208669_s_at	0.013223	gb:AF109873.1 /DEF=Homo sapiens retinoblastoma protein-associated protein mRNA, complete cds. /FEA=mRNA /PROD=retinoblastoma protein-associated protein /DB_XREF=gi:11415000 /UG=Hs.75847 CREBBPEP300 inhibitory protein 1 /FL=gb:AF109873.1 gb:AF274947.1 gb:AF274951.1 gb:AF092135.1 gb:NM_014335.1	AF109873		NP_055150
208671_at	1.26E-04	gb:AF164794.1 /DEF=Homo sapiens Diff33 protein homolog mRNA, complete cds. /FEA=mRNA /PROD=Diff33 protein homolog /DB_XREF=gi:8895090 /UG=Hs.146668 KIAA1253 protein /FL=gb:AF164794.1	AF164794		NP_065806
208674_x_at	1.26E-04	gb:BC002594.1 /DEF=Homo sapiens, dolichyl-diphosphooligosaccharide-protein glycosyltransferase, clone MGC:2191, mRNA, complete cds. /FEA=mRNA /PROD=dolichyl-diphosphooligosaccharide-proteinglycosyltransferase /DB_XREF=gi:12803530 /UG=Hs.34789 dolichyl-diphosphooligosaccharide-protein glycosyltransferase /FL=gb:BC002594.1 gb:D29643.1 gb:NM_005216.1	BC002594		NP_005207
208678_at	0.00161	gb:BC004443.1 /DEF=Homo sapiens, ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 31kD, clone MGC:4040, mRNA, complete cds. /FEA=mRNA /PROD=ATPase, H ⁺ transporting, lysosomal (vacuolarproton pump) 31kD /DB_XREF=gi:13325247 /UG=Hs.77805 ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 31kD /FL=gb:BC004443.1 gb:NM_001696.1	BC004443		NP_001687

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208679_s_at	7.58E-04	gb:AF279893.1 /DEF=Homo sapiens PNAS-139 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-139 /DB_XREF=gi:12751114 /UG=Hs.83583 actin related protein 23 complex, subunit 2 (34 kD) /FL=gb:BC000590.1 gb:AF279893.1	AF279893		NP_690601
208684_at	0.001116	gb:U24105.1 /DEF=Homo sapiens coatomer protein (COPA) mRNA, complete cds. /FEA=mRNA /GEN=COPA /PROD=coatomer protein /DB_XREF=gi:1638873 /UG=Hs.75887 coatomer protein complex, subunit alpha /FL=gb:U24105.1 gb:NM_004371.2	U24105		NP_004362
208685_x_at	0.001054	bromodomain containing 2	AA902767	Hs.75243	NP_005095
208689_s_at	0.033381	gb:BC003560.1 /DEF=Homo sapiens, ribophorin II, clone MGC:1817, mRNA, complete cds. /FEA=mRNA /PROD=ribophorin II /DB_XREF=gi:13097707 /UG=Hs.75722 ribophorin II /FL=gb:BC003560.1 gb:NM_002951.1	BC003560		NP_002942
208691_at	0.001116	gb:BC001188.1 /DEF=Homo sapiens, transferrin receptor (p90, CD71), clone MGC:3151, mRNA, complete cds. /FEA=mRNA /PROD=transferrin receptor (p90, CD71) /DB_XREF=gi:12654696 /UG=Hs.77356 transferrin receptor (p90, CD71) /FL=gb:BC001188.1 gb:M11507.1	BC001188		NP_003225
208692_at	0.013223	gb:U14990.1 /DEF=Human XP1PO ribosomal protein S3 (rpS3) mRNA, complete cds. /FEA=mRNA /GEN=rpS3 /PROD=ribosomal protein S3 /DB_XREF=gi:555940 /UG=Hs.252259 ribosomal protein S3 /FL=gb:BC003137.1 gb:NM_001005.1 gb:U14990.1 gb:U14991.1 gb:U14992.1	U14990		NP_000996

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208696_at	1.26E-04	gb:AF275798.1 /DEF=Homo sapiens PNAS-102 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-102 /DB_XREF=gi:10834761 /UG=Hs.1600 chaperonin containing TCP1, subunit 5 (epsilon) /FL=gb:AF275798.1	AF275798		NP_036205
208702_x_at	0.003171	Consensus includes gb:AI525212 /FEA=EST /DB_XREF=gi:4439347 /DB_XREF=est:pt1.1-2.A08.r /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2 /FL=gb:BC000373.1	BC000373		NP_001633
208703_s_at	0.026891	Consensus includes gb:BG427393 /FEA=EST /DB_XREF=gi:13333995 /DB_XREF=est:602499110F1 /CLONE=IMAGE:4612562 /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2 /FL=gb:BC000373.1	BC000373		NP_001633
208704_x_at	2.05E-04	gb:BC000373.1 /DEF=Homo sapiens, Similar to amyloid beta (A4) precursor-like protein 2; clone MGC:8371, mRNA, complete cds. /FEA=mRNA /PROD=Similar to amyloid beta (A4) precursor-like protein 2 /DB_XREF=gi:12653210 /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2 /FL=gb:BC000373.1	BC000373		NP_001633
208706_s_at	0.041055	Consensus includes gb:AK026933.1 /DEF=Homo sapiens cDNA: FLJ23280 fis, clone HEP07194. /FEA=mRNA /DB_XREF=gi:10439907 /UG=Hs.286236 eukaryotic translation initiation factor 5 /FL=gb:AL080102.1	AL080102		NP_001960

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208708_x_at	1.26E-04	gb:AL080102.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564N1916 (from clone DKFZp564N1916); complete cds. /FEA=mRNA /GEN=DKFZp564N1916 /PROD=hypothetical protein /DB_XREF=gi:5262526 /UG=Hs.286236 eukaryotic translation initiation factor 5 /FL=gb:AL080102.1	AL080102		NP_001960
208709_s_at	5.03E-04	gb:U64898.1 /DEF=Homo sapiens NRD convertase mRNA, complete cds. /FEA=mRNA /PROD=NRD convertase /DB_XREF=gi:2897866 /UG=Hs.4099 nardilysin (N-arginine dibasic convertase) /FL=gb:U64898.1	U64898		NP_002516
208713_at	0.013223	E1B-55kDa-associated protein 5	BF724216	Hs.155218	NP_653335
208716_s_at	0.001116	gb:AB020980.1 /DEF=Homo sapiens mRNA for putative membrane protein, complete cds. /FEA=mRNA /PROD=membrane protein /DB_XREF=gi:6467174 /UG=Hs.93832 putative membrane protein /FL=gb:BC000104.1 gb:AB020980.1	AB020980		NP_061899
208723_at	0.046906	gb:BC000350.1 /DEF=Homo sapiens, ubiquitin specific protease 11, clone MGC:8620, mRNA, complete cds. /FEA=mRNA /PROD=ubiquitin specific protease 11 /DB_XREF=gi:12653164 /UG=Hs.171501 ubiquitin specific protease 11 /FL=gb:BC000350.1 gb:U44839.1 gb:NM_004651.1	BC000350		NP_004642

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208724_s_at	1.26E-04	gb:BC000905.1 /DEF=Homo sapiens, RAB1, member RAS oncogene family, clone MGC:5233, mRNA, complete cds. /FEA=mRNA /PROD=RAB1, member RAS oncogene family /DB_XREF=gi:12654174 /UG=Hs.3642 RAB1, member RAS oncogene family /FL=gb:BC000905.1	BC000905		NP_004152
208726_s_at	0.02145	gb:BC000461.1 /DEF=Homo sapiens, eukaryotic translation initiation factor 2, subunit 2 (beta, 38kD), clone MGC:8508, mRNA, complete cds. /FEA=mRNA /PROD=eukaryotic translation initiation factor 2,subunit 2 (beta, 38kD) /DB_XREF=gi:12653382 /UG=Hs.12163 eukaryotic translation initiation factor 2, subunit 2 (beta, 38kD) /FL=gb:BC000461.1 gb:BC000934.2 gb:M29536.1 gb:NM_003908.1	BC000461		NP_003899
208728_s_at	3.26E-04	gb:BC003682.1 /DEF=Homo sapiens, cell division cycle 42 (GTP-binding protein, 25kD), clone MGC:5044, mRNA, complete cds. /FEA=mRNA /PROD=cell division cycle 42 (GTP-binding protein,25kD) /DB_XREF=gi:13277547 /UG=Hs.146409 cell division cycle 42 (GTP-binding protein, 25kD) /FL=gb:BC002711.1 gb:BC003682.1 gb:M57298.1 gb:NM_001791.1	BC003682		NP_426359

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208729_x_at	0.02145	gb:D83043.1 /DEF=Human HLA-B mRNA, allele A*2711, complete cds. /FEA=mRNA /GEN=HLA-B /DB_XREF=gi:1871135 /UG=Hs.77961 major histocompatibility complex, class I, B /FL=gb:U36492.1 gb:U35431.1 gb:L37881.1 gb:U31971.1 gb:L42345.1 gb:U50710.1 gb:L41925.1 gb:L41214.1 gb:U27608.1 gb:U35734.1 gb:U70528.1 gb:U70529.1 gb:U80945.1 gb:U64801.1 gb:U09864.1 gb:M77776.1 gb:M77775.1 gb:L04696.1 gb:D83043.1 gb:M12678.1 gb:L09736.1 gb:L09735.1 gb:M75138.1 gb:M84381.1 gb:M84382.1 gb:M84383.1 gb:M84384.1 gb:M84385.1 gb:M32318.1 gb:M32319.1 gb:M32320.1 gb:M81798.1 gb:L04695.1 gb:M84694.1 gb:U30936.1 gb:AF016641.1 gb:U36392.1 gb:AB008102.1 gb:U75533.1 gb:U57966.1 gb:AF026218.1 gb:L07743.1 gb:U01848.1 gb:U38800.1 gb:U40498.1 gb:U03027.1 gb:U03859.1 gb:L22028.1 gb:U04243.1 gb:U06862.1 gb:L32862.1	D83043		NP_005505
208730_x_at	0.004341	Consensus includes gb:AA535244 /FEA=EST /DB_XREF=gi:2279497 /DB_XREF=est:nf93a03.s1 /CLONE=IMAGE:927436 /UG=Hs.78305 RAB2, member RAS oncogene family /FL=gb:NM_002865.1 gb:M28213.1	NM_002865		NP_002856
208731_at	3.26E-04	Consensus includes gb:AU158062 /FEA=EST /DB_XREF=gi:11019583 /DB_XREF=est:AU158062 /CLONE=PLACE1010787 /UG=Hs.78305 RAB2, member RAS oncogene family /FL=gb:NM_002865.1 gb:M28213.1	NM_002865		NP_002856

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208732_at	3.26E-04	Consensus includes gb:AI743756 /FEA=EST /DB_XREF=gi:5112044 /DB_XREF=est:wg53e08.x1 /CLONE=IMAGE:2368838 /UG=Hs.78305 RAB2, member RAS oncogene family /FL=gb:NM_002865.1 gb:M28213.1	NM_002865		NP_002856
208734_x_at	0.010205	gb:M28213.1 /DEF=Homo sapiens GTP-binding protein (RAB2) mRNA, complete cds. /FEA=mRNA /GEN=RAB2 /PROD=GTP-binding protein /DB_XREF=gi:550061 /UG=Hs.78305 RAB2, member RAS oncogene family /FL=gb:NM_002865.1 gb:M28213.1	M28213		NP_002856
208736_at	1.26E-04	gb:AF004561.1 /DEF=Homo sapiens p21-Arc mRNA, complete cds. /FEA=mRNA /PROD=p21-Arc /DB_XREF=gi:2209346 /UG=Hs.6895 actin related protein 23 complex, subunit 3 (21 kD) /FL=gb:AF004561.1 gb:AF006086.1 gb:NM_005719.1	AF004561		NP_005710
208739_x_at	3.26E-04	gb:L76416.1 /DEF=Homo sapiens MIF2 suppressor (HSMT3) mRNA, complete cds. /FEA=mRNA /GEN=HSMT3 /PROD=MIF2 suppressor /DB_XREF=gi:5566605 /UG=Hs.180139 SMT3 (suppressor of mif two 3, yeast) homolog 2 /FL=gb:L76416.1 gb:NM_006937.1	L76416		NP_008868
208741_at	0.004341	Consensus includes gb:AW274856 /FEA=EST /DB_XREF=gi:6661886 /DB_XREF=est:xm61g03.x1 /CLONE=IMAGE:2688724 /UG=Hs.23964 sin3-associated polypeptide, 18kD /FL=gb:NM_005870.2 gb:U96915.1 gb:AF153608.1 gb:U78303.1	NM_005870		NP_005861

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208742_s_at	1.26E-04	gb:U78303.1 /DEF=Human 2HOR0202 mRNA, complete cds. /FEA=mRNA /PROD=2HOR0202 /DB_XREF=gi:6648546 /UG=Hs.23964 sin3-associated polypeptide, 18kD /FL=gb:NM_005870.2 gb:U96915.1 gb:AF153608.1 gb:U78303.1	U78303		NP_005861
208743_s_at	7.58E-04	gb:BC001359.1 /DEF=Homo sapiens, tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, beta polypeptide, clone MGC:1580, mRNA, complete cds. /FEA=mRNA /PROD=tyrosine 3-monooxygenasetryptophan5-monooxygenase activation protein, beta polypeptide /DB_XREF=gi:12655022 /UG=Hs.279920 tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, beta polypeptide /FL=gb:BC001359.1	BC001359		NP_647539
208746_x_at	5.03E-04	gb:AF070655.1 /DEF=Homo sapiens F1F0-type ATP synthase subunit g mRNA, complete cds. /FEA=mRNA /PROD=F1F0-type ATP synthase subunit g /DB_XREF=gi:4454685 /UG=Hs.107476 ATP synthase, H ⁺ transporting, mitochondrial F1F0, subunit g /FL=gb:AF070655.1	AF070655		NP_006467
208749_x_at	2.05E-04	gb:AF085357.1 /DEF=Homo sapiens flotillin mRNA, complete cds. /FEA=mRNA /PROD=flotillin /DB_XREF=gi:5114048 /UG=Hs.179986 flotillin 1 /FL=gb:BC001146.1 gb:AF085357.1 gb:AF089750.2 gb:NM_005803.2	AF085357		NP_005794

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208756_at	0.010205	gb:U36764.1 /DEF=Human TGF-beta receptor interacting protein 1 mRNA, complete cds. /FEA=mRNA /PROD=TGF-beta receptor interacting protein 1 /DB_XREF=gi:1036804 /UG=Hs.192023 eukaryotic translation initiation factor 3, subunit 2 (beta, 36kD) /FL=gb:U36764.1 gb:BC000413.1 gb:BC003140.1 gb:U39067.1 gb:NM_003757.1 gb:AF116697.1	U36764		NP_003748
208757_at	0.035171	gb:BC001123.1 /DEF=Homo sapiens, Similar to gp25L2 protein, clone MGC:2142, mRNA, complete cds. /FEA=mRNA /PROD=Similar to gp25L2 protein /DB_XREF=gi:12654574 /UG=Hs.279929 gp25L2 protein /FL=gb:BC001123.1	BC001123		NP_059980
208759_at	0.003171	gb:AF240468.1 /DEF=Homo sapiens nicastrin mRNA, complete cds. /FEA=mRNA /PROD=nicastrin /DB_XREF=gi:9992877 /UG=Hs.4788 nicastrin /FL=gb:AF240468.1	AF240468		NP_056146
208761_s_at	1.26E-04	gb:U67122.1 /DEF=Human ubiquitin-related protein SUMO-1 mRNA, complete cds. /FEA=mRNA /PROD=SUMO-1 /DB_XREF=gi:1762972 /UG=Hs.81424 ubiquitin-like 1 (sentrin) /FL=gb:U61397.1 gb:U38784.1 gb:U72722.1 gb:U67122.1 gb:U83117.1 gb:NM_003352.1	U67122		NP_003343
208764_s_at	0.003171	gb:D13119.1 /DEF=Homo sapiens P2 mRNA for ATP synthase subunit c, complete cds. /FEA=mRNA /GEN=P2 /PROD=ATP synthase subunit c precursor /DB_XREF=gi:285909 /UG=Hs.89399 ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2 /FL=gb:D13119.1	D13119		NP_005167

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208765_s_at	0.010205	Consensus includes gb:NM_005826.1 /DEF=Homo sapiens heterogeneous nuclear ribonucleoprotein R (HNRPR), mRNA. /FEA=CDS /GEN=HNRPR /PROD=heterogeneous nuclear ribonucleoprotein R /DB_XREF=gi:5031754 /UG=Hs.15265 heterogeneous nuclear ribonucleoprotein R /FL=gb:BC001449.1 gb:AF000364.1 gb:NM_005826.1	NM_005826		NP_005817
208767_s_at	0.001116	putative integral membrane transporter	AW149681	Hs.296398	NP_060877
208770_s_at	2.05E-04	gb:BC005057.1 /DEF=Homo sapiens, eukaryotic translation initiation factor 4E binding protein 2, clone MGC:12944, mRNA, complete cds. /FEA=mRNA /PROD=eukaryotic translation initiation factor 4Ebinding protein 2 /DB_XREF=gi:13477190 /UG=Hs.278712 eukaryotic translation initiation factor 4E binding protein 2 /FL=gb:BC005057.1 gb:NM_004096.1 gb:L36056.1	BC005057		NP_004087
208771_s_at	1.26E-04	gb:J02959.1 /DEF=Human leukotriene A-4 hydrolase mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:187174 /UG=Hs.81118 leukotriene A4 hydrolase /FL=gb:J03459.1 gb:J02959.1 gb:NM_000895.1	J02959		NP_000886
208773_s_at	0.016934	gb:AL136943.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586G1024 (from clone DKFZp586G1024); complete cds. /FEA=mRNA /GEN=DKFZp586G1024 /PROD=hypothetical protein /DB_XREF=gi:12053380 /UG=Hs.301226 KIAA1085 protein /FL=gb:AL136943.1	AL136943		NP_078944

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208780_x_at	1.26E-04	gb:AF154847.1 /DEF=Homo sapiens 33 kDa Vamp-associated protein (VAMP) mRNA, complete cds. /FEA=mRNA /GEN=VAMP /PROD=33 kDa Vamp-associated protein /DB_XREF=gi:8099349 /UG=Hs.9006 VAMP (vesicle-associated membrane protein)-associated protein A (33kD) /FL=gb:BC002992.1 gb:AF057358.1 gb:AF044670.1 gb:AF086627.1 gb:NM_003574.1 gb:AF154847.1	AF154847		NP_003565
208781_x_at	1.26E-04	gb:AF062483.1 /DEF=Homo sapiens SDP3 mRNA, complete cds. /FEA=mRNA /PROD=SDP3 /DB_XREF=gi:3126978 /UG=Hs.12102 sorting nexin 3 /FL=gb:AF062483.1 gb:AF034546.1 gb:NM_003795.1	AF062483		NP_690041
208783_s_at	0.016934	membrane cofactor protein (CD46, trophoblast-lymphocyte cross-reactive antigen)	AL570661	Hs.83532	NP_758871
208785_s_at	2.05E-04	ESTs, Highly similar to MPL3_HUMAN Microtubule-associated proteins 1A/1B light chain 3 (MAP1A/MAP1B LC3) [H.sapiens]	BE893893	Hs.356061	
208786_s_at	1.26E-04	gb:AF183417.1 /DEF=Homo sapiens microtubule-associated proteins 1A1B light chain 3 mRNA, complete cds. /FEA=mRNA /PROD=microtubule-associated proteins 1A1B lightchain 3 /DB_XREF=gi:9963772 /UG=Hs.121849 microtubule-associated proteins 1A1B light chain 3 /FL=gb:AF303888.1 gb:NM_022818.2 gb:AF183417.1	AF183417		NP_073729

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208788_at	0.007779	gb:AL136939.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586B1824 (from clone DKFZp586B1824); complete cds. /FEA=mRNA /GEN=DKFZp586B1824 /PROD=hypothetical protein /DB_XREF=gi:12053372 /UG=Hs.250175 homolog of yeast long chain polyunsaturated fatty acid elongation enzyme 2 /FL=gb:NM_021814.1 gb:AL136939.1 gb:AF111849.1 gb:AF231981.1	AL136939		NP_068586
208791_at	0.003026	gb:M25915.1 /DEF=Human complement cytolysis inhibitor (CLI) mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:180619 /UG=Hs.75106 clusterin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J) /FL=gb:J02908.1 gb:M25915.1 gb:M64722.1 gb:NM_001831.1	M25915		NP_001822
208792_s_at	3.26E-04	gb:M25915.1 /DEF=Human complement cytolysis inhibitor (CLI) mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:180619 /UG=Hs.75106 clusterin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J) /FL=gb:J02908.1 gb:M25915.1 gb:M64722.1 gb:NM_001831.1	M25915		NP_001822
208797_s_at	0.018206	golgin-67	AI829170	Hs.182982	NP_851422

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208799_at	1.26E-04	gb:BC004146.1 /DEF=Homo sapiens, proteasome (prosome, macropain) subunit, beta type, 5, clone MGC:2175, mRNA, complete cds. /FEA=mRNA /PROD=proteasome (prosome, macropain) subunit, betatype, 5 /DB_XREF=gi:13278740 /UG=Hs.78596 proteasome (prosome, macropain) subunit, beta type, 5 /FL=gb:BC004146.1 gb:NM_002797.1 gb:D29011.1	BC004146		
208800_at	0.02145	Consensus includes gb:AV702627 /FEA=EST /DB_XREF=gi:10718957 /DB_XREF=est:AV702627 /CLONE=ADBAHC04 /UG=Hs.237825 signal recognition particle 72kD /FL=gb:AF069765.1 gb:AF077019.1 gb:NM_006947.1	NM_006947		NP_008878
208805_at	7.58E-04	gb:BC002979.1 /DEF=Homo sapiens, proteasome (prosome, macropain) subunit, alpha type 6, clone MGC:2333, mRNA, complete cds. /FEA=mRNA /PROD=proteasome (prosome, macropain) subunit, alphas type 6 /DB_XREF=gi:12804240 /UG=Hs.74077 proteasome (prosome, macropain) subunit, alpha type, 6 /FL=gb:BC002979.1	BC002979		NP_002782
208807_s_at	0.016426	gb:U91543.1 /DEF=Homo sapiens zinc-finger helicase (hZFH) mRNA, complete cds. /FEA=mRNA /GEN=hZFH /PROD=zinc-finger helicase /DB_XREF=gi:3298561 /UG=Hs.25601 chromodomain helicase DNA binding protein 3 /FL=gb:U91543.1	U91543		NP_001263

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208808_s_at	1.26E-04	gb:BC000903.1 /DEF=Homo sapiens, high-mobility group (nonhistone chromosomal) protein 2, clone MGC:5234, mRNA, complete cds. /FEA=mRNA /PROD=high-mobility group (nonhistone chromosomal)protein 2 /DB_XREF=gi:12654170 /UG=Hs.80684 high-mobility group (nonhistone chromosomal) protein 2 /FL=gb:BC000903.1 gb:BC001063.1	BC000903		NP_002120
208809_s_at	0.005852	gb:AL136632.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G182 (from clone DKFZp564G182); complete cds. /FEA=mRNA /GEN=DKFZp564G182 /PROD=hypothetical protein /DB_XREF=gi:12052789 /UG=Hs.173685 hypothetical protein FLJ12619 /FL=gb:AL136632.1	AL136632		NP_112201
208810_at	0.013223	gb:AF080569.1 /DEF=Homo sapiens DnaJ-like 2 protein (HSJ2) mRNA, complete cds. /FEA=mRNA /GEN=HSJ2 /PROD=DnaJ-like 2 protein /DB_XREF=gi:4322314 /UG=Hs.181195 DnaJ (Hsp40) homolog, subfamily B, member 6 /FL=gb:AL136707.1 gb:BC000177.2 gb:AB014888.1 gb:AF080569.1 gb:NM_005494.1 gb:AF075601.1 gb:AF060703.1 gb:AB015799.1	AF080569		NP_490647
208811_s_at	0.010205	gb:AF080569.1 /DEF=Homo sapiens DnaJ-like 2 protein (HSJ2) mRNA, complete cds. /FEA=mRNA /GEN=HSJ2 /PROD=DnaJ-like 2 protein /DB_XREF=gi:4322314 /UG=Hs.181195 DnaJ (Hsp40) homolog, subfamily B, member 6 /FL=gb:AL136707.1 gb:BC000177.2 gb:AB014888.1 gb:AF080569.1 gb:NM_005494.1 gb:AF075601.1 gb:AF060703.1 gb:AB015799.1	AF080569		NP_490647

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208812_x_at	0.026891	gb:BC004489.1 /DEF=Homo sapiens, major histocompatibility complex, class I, C, clone MGC:11039, mRNA, complete cds. /FEA=mRNA /PROD=major histocompatibility complex, class I, C /DB_XREF=gi:13325360 /UG=Hs.277477 major histocompatibility complex, class I, C /FL=gb:NM_002117.1 gb:M99388.1 gb:U06487.1 gb:BC002463.1 gb:BC004489.1 gb:D64145.1 gb:D38526.1 gb:D49552.1 gb:D49819.1 gb:M24097.1 gb:M84171.1 gb:M84172.1 gb:M84173.1 gb:M84174.1 gb:M26429.1 gb:M26430.1 gb:M26431.1 gb:U41420.1 gb:U41386.1 gb:D50852.1 gb:D50853.1 gb:D50854.1 gb:D83031.1 gb:U57028.1 gb:U06695.1 gb:U06696.1 gb:M99389.1 gb:M99390.1 gb:M28160.1 gb:U09853.1 gb:AF168611.1 gb:L38251.1 gb:D31817.1	BC004489		NP_002108
208815_x_at	0.010205	gb:AB023420.1 /DEF=Homo sapiens mRNA for heat shock protein apg-2, complete cds. /FEA=mRNA /GEN=apg-2 /PROD=apg-2 /DB_XREF=gi:4579908 /UG=Hs.90093 heat shock 70kD protein 4 /FL=gb:AB023420.1	AB023420		
208818_s_at	0.02145	gb:BC000419.1 /DEF=Homo sapiens, catechol-O-methyltransferase, clone MGC:8663, mRNA, complete cds. /FEA=mRNA /PROD=catechol-O-methyltransferase /DB_XREF=gi:12653300 /UG=Hs.240013 catechol-O-methyltransferase /FL=gb:BC000419.1 gb:M58525.1 gb:M65212.1 gb:NM_007310.1 gb:NM_000754.2	BC000419		NP_009294

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208825_x_at	0.00161	gb:U43701.1 /DEF=Human ribosomal protein L23a mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein L23a /DB_XREF=gi:1399085 /UG=Hs.184776 ribosomal protein L23a /FL=gb:U43701.1	U43701		NP_000975
208834_x_at	0.00161	gb:BC001865.1 /DEF=Homo sapiens, Similar to cadherin 1, type 1, E-cadherin (epithelial), clone MGC:1151, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cadherin 1, type 1, E-cadherin(epithelial) /DB_XREF=gi:12804838 /UG=Hs.194657 cadherin 1, type 1, E-cadherin (epithelial) /FL=gb:BC001865.1	BC001865		
208835_s_at	1.26E-04	cisplatin resistance-associated overexpressed protein	AW089673	Hs.3688	NP_057508
208857_s_at	1.26E-04	gb:M93008.1 /DEF=Human L-isoaspartylD-aspartyl protein carboxyl methyltransferase isozyme II mRNA, complete cds. /FEA=mRNA /PROD=L-isoaspartylD-aspartyl protein carboxylmethyltransferase /DB_XREF=gi:180636 /UG=Hs.79137 protein-L-isoaspartate (D-aspartate) O-methyltransferase /FL=gb:D25546.1 gb:M93008.1 gb:D13892.1	M93008		NP_005380
208859_s_at	0.018693	alpha thalassemia/mental retardation syndrome X-linked (RAD54 homolog, <i>S. cerevisiae</i>)	AI650257	Hs.96264	NP_612115
208864_s_at	2.05E-04	gb:AF313911.1 /DEF=Homo sapiens thioredoxin mRNA, complete cds. /FEA=mRNA /PROD=thioredoxin /DB_XREF=gi:11345419 /UG=Hs.76136 thioredoxin /FL=gb:AF313911.1 gb:BC003377.1 gb:J04026.1 gb:NM_003329.1 gb:AF276919.1 gb:AY004872.1	AF313911		NP_003320
208865_at	2.05E-04	casein kinase 1, alpha 1	BG534245	Hs.283738	NP_001883

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208867_s_at	0.033381	gb:AF119911.1 /DEF=Homo sapiens PRO2975 mRNA, complete cds. /FEA=mRNA /PROD=PRO2975 /DB_XREF=gi:7770258 /UG=Hs.144477 hypothetical protein PRO2975 /FL=gb:AF119911.1	AF119911		
208870_x_at	0.02145	gb:BC000931.2 /DEF=Homo sapiens, ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1, clone MGC:5380, mRNA, complete cds. /FEA=mRNA /PROD=ATP synthase, H+ transporting, mitochondrial F1complex, gamma polypeptide 1 /DB_XREF=gi:13111822 /UG=Hs.155433 ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1 /FL=gb:BC000470.1 gb:BC000931.2 gb:D16562.1	BC000931		NP_005165
208872_s_at	1.26E-04	likely ortholog of mouse deleted in polyposis 1	AA814140	Hs.178112	NP_005660
208873_s_at	0.005852	gb:BC000232.1 /DEF=Homo sapiens, Similar to deleted in polyposis 1, clone MGC:2267, mRNA, complete cds. /FEA=mRNA /PROD=Similar to deleted in polyposis 1 /DB_XREF=gi:12652946 /UG=Hs.178112 DNA segment, single copy probe LNS-CAILNS-CAII (deleted in polyposis /FL=gb:BC000232.1	BC000232		NP_005660
208875_s_at	0.00161	Consensus includes gb:BF796470 /FEA=EST /DB_XREF=gi:12101524 /DB_XREF=est:602259926F1 /CLONE=IMAGE:4342999 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF092132.1	AF092132		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208876_s_at	0.013223	Consensus includes gb:AI076186 /FEA=EST /DB_XREF=gi:3405364 /DB_XREF=est:oz01g01.x1 /CLONE=IMAGE:1674096 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF092132.1	AF092132		
208877_at	0.010205	Consensus includes gb:W74494 /FEA=EST /DB_XREF=gi:1384781 /DB_XREF=est:zd75f10.s1 /CLONE=IMAGE:346507 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF092132.1	AF092132		
208878_s_at	0.001116	gb:AF092132.1 /DEF=Homo sapiens PAK2 mRNA, complete cds. /FEA=mRNA /PROD=PAK2 /DB_XREF=gi:5138913 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF092132.1	AF092132		
208880_s_at	0.009689	gb:AB019219.1 /DEF=Homo sapiens mRNA, complete cds, similar to yeast pre-mRNA splicing factors, Prp1Zer1 and Prp6. /FEA=mRNA /DB_XREF=gi:4164165 /UG=Hs.31334 putative mitochondrial outer membrane protein import receptor /FL=gb:BC001666.1 gb:AF026031.1 gb:AB019219.1 gb:NM_012469.1 gb:AF221842.1	AB019219		NP_036601
208881_x_at	1.26E-04	gb:BC005247.1 /DEF=Homo sapiens, isopentenyl-diphosphate delta isomerase, clone MGC:12281, mRNA, complete cds. /FEA=mRNA /PROD=isopentenyl-diphosphate delta isomerase /DB_XREF=gi:13528899 /UG=Hs.76038 isopentenyl-diphosphate delta isomerase /FL=gb:BC005247.1	BC005247		NP_004499
208882_s_at	3.26E-04	progesterone induced protein	U69567	Hs.278428	NP_056986

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208884_s_at	1.26E-04	gb:AF006010.1 /DEF=Homo sapiens progesterone induced protein (EDD) mRNA, complete cds. /FEA=mRNA /GEN=EDD /PROD=progesterone induced protein /DB_XREF=gi:4101694 /UG=Hs.278428 progesterone induced protein /FL=gb:NM_015902.2 gb:AF006010.1 gb:U95000.1	AF006010		NP_056986
208886_at	1.28E-04	gb:BC000145.1 /DEF=Homo sapiens, H1 histone family, member 0, clone MGC:5241, mRNA, complete cds. /FEA=mRNA /PROD=H1 histone family, member 0 /DB_XREF=gi:12652786 /UG=Hs.226117 H1 histone family, member 0 /FL=gb:BC000145.1 gb:NM_005318.1	BC000145		NP_005309
208887_at	0.005852	gb:BC000733.1 /DEF=Homo sapiens, eukaryotic translation initiation factor 3, subunit 4 (delta, 44kD), clone MGC:2053, mRNA, complete cds. /FEA=mRNA /PROD=eukaryotic translation initiation factor 3, subunit 4 (delta, 44kD) /DB_XREF=gi:12653882 /UG=Hs.28081 eukaryotic translation initiation factor 3, subunit 4 (delta, 44kD) /FL=gb:AF094850.1 gb:BC000733.1 gb:AF020833.1 gb:U96074.1 gb:NM_003755.1	BC000733		NP_003746
208894_at	0.02145	gb:M60334.1 /DEF=Human MHC class II HLA-DR-alpha mRNA, complete cds. /FEA=mRNA /GEN=HLA-DRA /PROD=cell surface glycoprotein /DB_XREF=gi:188255 /UG=Hs.76807 major histocompatibility complex, class II, DR alpha /FL=gb:M60334.1 gb:NM_019111.1	M60334		NP_061984

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208895_s_at	0.00161	Consensus includes gb:BG530850 /FEA=EST /DB_XREF=gi:13522387 /DB_XREF=est:602559887F1 /CLONE=IMAGE:4698044 /UG=Hs.100555 DEADH (Asp-Glu-Ala-AspHis) box polypeptide 18 (Myc-regulated) /FL=gb:BC001238.1 gb:BC003360.1	BC003360		NP_006764
208898_at	2.05E-04	gb:AF077614.1 /DEF=Homo sapiens vacuolar ATP synthase subunit D homolog (VATD) mRNA, complete cds. /FEA=mRNA /GEN=VATD /PROD=vacuolar ATP synthase subunit D homolog /DB_XREF=gi:11999089 /UG=Hs.272630 vacuolar proton pump delta polypeptide /FL=gb:AF104629.1 gb:AF077614.1 gb:BC001411.1 gb:AF145316.1 gb:AF100741.1 gb:NM_015994.1	AF077614		NP_057078
208899_x_at	1.26E-04	gb:AF100741.1 /DEF=Homo sapiens vacuolar H-ATPase subunit D mRNA, complete cds. /FEA=mRNA /PROD=vacuolar H-ATPase subunit D /DB_XREF=gi:5138992 /UG=Hs.272630 vacuolar proton pump delta polypeptide /FL=gb:AF104629.1 gb:AF077614.1 gb:BC001411.1 gb:AF145316.1 gb:AF100741.1 gb:NM_015994.1	AF100741		NP_057078
208900_s_at	0.003539	topoisomerase (DNA) I	AW025108	Hs.317	NP_003277
208904_s_at	0.041055	gb:BC000354.1 /DEF=Homo sapiens, ribosomal protein S28, clone MGC:8658, mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein S28 /DB_XREF=gi:12653172 /UG=Hs.153177 ribosomal protein S28 /FL=gb:BC000354.1	BC000354		NP_001022

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208908_s_at	0.003171	gb:AF327443.1 /DEF=Homo sapiens calpastatin mRNA, complete cds. /FEA=mRNA /PROD=calpastatin /DB_XREF=gi:12056961 /UG=Hs.279607 calpastatin /FL=gb:AF327443.1 gb:U26724.2	AF327443		NP_775085
208909_at	1.26E-04	gb:BC000649.1 /DEF=Homo sapiens, ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1, clone MGC:1362, mRNA, complete cds. /FEA=mRNA /PROD=ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1 /DB_XREF=gi:12653726 /UG=Hs.3712 ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1 /FL=gb:BC000649.1 gb:NM_006003.1	BC000649		NP_005994
208911_s_at	0.007258	gb:M34055.1 /DEF=Human pyruvate dehydrogenase E1-beta subunit mRNA, complete cds. /FEA=mRNA /GEN=PDHB /DB_XREF=gi:190791 /UG=Hs.979 pyruvate dehydrogenase (lipoamide) beta /FL=gb:BC000439.1 gb:BC001924.1 gb:J03576.1 gb:M34479.1 gb:M54788.1 gb:M34055.1 gb:NM_000925.1	M34055		NP_000916
208913_at	0.010205	Consensus includes gb:AA868560 /FEA=EST /DB_XREF=gi:2964005 /DB_XREF=est:ak43g11.s1 /CLONE=IMAGE:1408772 /UG=Hs.155546 KIAA1080 protein; Golgi-associated, gamma-adaptin ear containing, ARF-binding protein 2 /FL=gb:AF190863.1 gb:AF233522.1 gb:AF165531.1 gb:NM_015044.1	NM_015044		NP_619581

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208918_s_at	0.013223	Consensus includes gb:AI334128 /FEA=EST /DB_XREF=gi:4070687 /DB_XREF=est:qq07e12.x1 /CLONE=IMAGE:1931854 /UG=Hs.220324 hypothetical protein FLJ13052 /FL=gb:BC001709.1	BC001709		NP_075394
208919_s_at	5.03E-04	gb:BC001709.1 /DEF=Homo sapiens, Similar to hypothetical protein FLJ13052, clone MGC:1900, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein FLJ13052 /DB_XREF=gi:12804578 /UG=Hs.220324 hypothetical protein FLJ13052 /FL=gb:BC001709.1	BC001709		NP_075394
208921_s_at	2.05E-04	gb:L12387.1 /DEF=Human sorcin (SRI) mRNA, complete cds. /FEA=mRNA /GEN=SRI /PROD=sorcin /DB_XREF=gi:459835 /UG=Hs.300741 sorcin /FL=gb:M32886.1 gb:NM_003130.1 gb:L12387.1	L12387		NP_003121
208923_at	3.26E-04	gb:BC005097.1 /DEF=Homo sapiens, Similar to selective hybridizing clone, clone MGC:13167, mRNA, complete cds. /FEA=mRNA /PROD=Similar to selective hybridizing clone /DB_XREF=gi:13477256 /UG=Hs.77257 KIAA0068 protein /FL=gb:BC005097.1	BC005097		NP_055423
208924_at	3.26E-04	gb:AB024703.1 /DEF=Homo sapiens mRNA for Sid1669p, complete cds. /FEA=mRNA /GEN=Sid1669 /PROD=Sid1669p /DB_XREF=gi:5931544 /UG=Hs.96334 ring finger protein 11 /FL=gb:AF151881.1 gb:AB024703.1 gb:NM_014372.1	AB024703		NP_055187

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208926_at	8.34E-04	gb:U84246.1 /DEF=Homo sapiens lysosomal sialidase mRNA, complete cds. /FEA=mRNA /PROD=lysosomal sialidase /DB_XREF=gi:4099140 /UG=Hs.118721 sialidase 1 (lysosomal sialidase) /FL=gb:BC000722.1 gb:AF040958.1 gb:U84246.1 gb:NM_000434.1	U84246		NP_000425
208929_x_at	0.026891	gb:BC004954.1 /DEF=Homo sapiens, clone MGC:10897, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10897) /DB_XREF=gi:13436331 /UG=Hs.180842 ribosomal protein L13 /FL=gb:BC000851.1 gb:BC004954.1 gb:NM_000977.1	BC004954		NP_150254
208932_at	0.037567	gb:BC001416.1 /DEF=Homo sapiens, protein phosphatase 4 (formerly X), catalytic subunit, clone MGC:1882, mRNA, complete cds. /FEA=mRNA /PROD=protein phosphatase 4 (formerly X), catalytic subunit /DB_XREF=gi:12655126 /UG=Hs.2903 protein phosphatase 4 (formerly X), catalytic subunit /FL=gb:BC001416.1 gb:AF097996.1 gb:NM_002720.1	BC001416		NP_002711
208933_s_at	0.02145	Fc fragment of IgG, low affinity IIb, receptor for (CD32)	AI659005	Hs.82316	NP_060542
208935_s_at	0.004341	gb:L78132.1 /DEF=Human prostate carcinoma tumor antigen (pcta-1) mRNA, complete cds. /FEA=mRNA /GEN=pcta-1 /PROD=prostate carcinoma tumor antigen /DB_XREF=gi:1932711 /UG=Hs.4082 lectin, galactoside-binding, soluble, 8 (galectin 8) /FL=gb:AF342815.1 gb:L78132.1 gb:AF074000.1 gb:NM_006499.1	L78132		NP_006490

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208943_s_at	0.001116	gb:U93239.1 /DEF=Human Sec62 (Sec62) mRNA, complete cds. /FEA=mRNA /GEN=Sec62 /PROD=Sec62 /DB_XREF=gi:1928972 /UG=Hs.8146 translocation protein 1 /FL=gb:D87127.1 gb:U93239.1 gb:NM_003262.1	U93239		NP_003253
208946_s_at	7.58E-04	gb:AF139131.1 /DEF=Homo sapiens beclin 1 (BECN1) mRNA, complete cds. /FEA=mRNA /GEN=BECN1 /PROD=beclin 1 /DB_XREF=gi:4680380 /UG=Hs.12272 beclin 1 (coiled-coil, myosin-like BCL2-interacting protein) /FL=gb:AF077301.1 gb:NM_003766.1 gb:AF139131.1	AF139131		NP_003757
208948_s_at	2.05E-04	gb:BC000830.1 /DEF=Homo sapiens, clone MGC:4921, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:4921) /DB_XREF=gi:12654050 /UG=Hs.6113 staufer (Drosophila, RNA-binding protein) /FL=gb:BC000830.1 gb:AF061938.1 gb:NM_017452.1	BC000830		NP_059348
208949_s_at	2.05E-04	gb:BC001120.1 /DEF=Homo sapiens, lectin, galactoside-binding, soluble, 3 (galectin 3), clone MGC:2058, mRNA, complete cds. /FEA=mRNA /PROD=lectin, galactoside-binding, soluble, 3(galectin 3) /DB_XREF=gi:12654570 /UG=Hs.621 lectin, galactoside-binding, soluble, 3 (galectin 3) /FL=gb:M35368.1 gb:BC001120.1 gb:M57710.1 gb:M36682.1 gb:AB006780.1 gb:NM_002306.1	BC001120		NP_002297

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208954_s_at	0.012367	gb:BC003381.1 /DEF=Homo sapiens, clone MGC:5353, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:5353) /DB_XREF=gi:13097242 /UG=Hs.78851 KIAA0217 protein /FL=gb:BC003381.1	BC003381		
208955_at	0.00228	gb:AB049113.1 /DEF=Homo sapiens DUT mRNA for dUTP pyrophosphatase, complete cds. /FEA=mRNA /GEN=DUT /PROD=dUTP pyrophosphatase /DB_XREF=gi:10257384 /UG=Hs.82113 dUTP pyrophosphatase /FL=gb:AB049113.1 gb:U31930.1 gb:U62891.1 gb:M89913.1 gb:NM_001948.1	AB049113		NP_001939
208959_s_at	0.004341	gb:BC005374.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 1110001E24 gene, clone MGC:12490, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 1110001E24 gene /DB_XREF=gi:13529223 /UG=Hs.154023 KIAA0573 protein /FL=gb:BC005374.1	BC005374		
208964_s_at	0.027243	gb:AL512760.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762M2311 (from clone DKFZp762M2311); complete cds. /FEA=mRNA /GEN=DKFZp762M2311 /PROD=hypothetical protein /DB_XREF=gi:12224983 /UG=Hs.132898 fatty acid desaturase 1 /FL=gb:AF084558.1 gb:NM_013402.2 gb:AL512760.1 gb:AF199596.1 gb:AF226273.1	AL512760		NP_037534
208966_x_at	0.02145	gb:AF208043.1 /DEF=Homo sapiens IFI16b (IFI16b) mRNA, complete cds. /FEA=mRNA /GEN=IFI16b /PROD=IFI16b /DB_XREF=gi:6644296 /UG=Hs.155530 interferon, gamma-inducible protein 16 /FL=gb:AF208043.1	AF208043		NP_005522

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208967_s_at	0.007779	gb:U39945.1 /DEF=Human adenylate kinase 2 (adk2) mRNA, complete cds. /FEA=mRNA /GEN=adk2 /PROD=adenylate kinase 2 /DB_XREF=gi:1209686 /UG=Hs.171811 adenylate kinase 2 /FL=gb:U39945.1 gb:U84371.1 gb:NM_001625.1	U39945		NP_751949
208970_s_at	7.58E-04	gb:M14016.1 /DEF=Human uroporphyrinogen decarboxylase mRNA, complete cds. /FEA=mRNA /GEN=UROD /DB_XREF=gi:340180 /UG=Hs.78601 uroporphyrinogen decarboxylase /FL=gb:BC001778.1 gb:M14016.1 gb:AF104421.1 gb:AF104422.1 gb:AF104423.1 gb:AF104424.1 gb:AF104425.1 gb:AF104426.1 gb:AF104427.1 gb:AF104428.1 gb:AF104429.1 gb:AF104430.1 gb:AF104431.1 gb:AF104432.1 gb:AF104433.1 gb:AF104434.1 gb:AF104435.1 gb:AF104436.1 gb:AF104437.1 gb:AF104438.1 gb:AF104439.1 gb:AF104440.1 gb:NM_000374.2	M14016		NP_000365
208977_x_at	0.00161	gb:BC004188.1 /DEF=Homo sapiens, tubulin, beta, 2, clone MGC:2826, mRNA, complete cds. /FEA=mRNA /PROD=tubulin, beta, 2 /DB_XREF=gi:13278848 /UG=Hs.251653 tubulin, beta, 2 /FL=gb:BC002783.1 gb:BC002885.1 gb:BC001911.1 gb:BC004188.1 gb:NM_006088.1	BC004188		NP_006079

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208979_at	1.26E-04	gb:AF128458.1 /DEF=Homo sapiens nuclear receptor coactivator RAP250 mRNA, complete cds. /FEA=mRNA /PROD=nuclear receptor coactivator RAP250 /DB_XREF=gi:7140573 /UG=Hs.159613 thyroid hormone receptor binding protein /FL=gb:AF177388.1 gb:AF208227.1 gb:AF128458.1 gb:NM_014071.1 gb:AF171667.1 gb:AF245115.1	AF128458		NP_054790
208981_at	1.26E-04	platelet/endothelial cell adhesion molecule (CD31 antigen)	AW574504	Hs.78146	NP_000433
208982_at	7.58E-04	platelet/endothelial cell adhesion molecule (CD31 antigen)	AW574504	Hs.78146	NP_000433
208988_at	0.033381	Consensus includes gb:BE675843 /FEA=EST /DB_XREF=gi:10036384 /DB_XREF=est:7f17b04.x1 /CLONE=IMAGE:3294895 /UG=Hs.219614 f-box and leucine rich repeat protein 11 /FL=gb:AF179221.1	AK024505		NP_036440
208989_s_at	0.003529	gb:AF179221.1 /DEF=Homo sapiens F-box protein Lilina (LILINA) mRNA, complete cds. /FEA=mRNA /GEN=LILINA /PROD=F-box protein Lilina /DB_XREF=gi:5917729 /UG=Hs.219614 f-box and leucine rich repeat protein 11 /FL=gb:AF179221.1	AF179221		NP_036440
208991_at	0.010205	signal transducer and activator of transcription 3 (acute-phase response factor)	AA634272	Hs.321677	
208993_s_at	6.68E-04	Consensus includes gb:AW340788 /FEA=EST /DB_XREF=gi:6837414 /DB_XREF=est:ha12h05.x1 /CLONE=IMAGE:2873529 /UG=Hs.77965 peptidyl-prolyl isomerase G (cyclophilin G) /FL=gb:U40763.1 gb:NM_004792.1	NM_004792		NP_004783

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
208994_s_at	7.58E-04	Consensus includes gb:A1638762 /FEA=EST /DB_XREF=gi:4690996 /DB_XREF=est:tt32a04.x1 /CLONE=IMAGE:2242446 /UG=Hs.77965 peptidyl-prolyl isomerase G (cyclophilin G) /FL=gb:U40763.1 gb:NM_004792.1	NM_004792		NP_004783
209003_at	0.007779	gb:AF070548.1 /DEF=Homo sapiens clone 24408 2-oxoglutarate carrier protein mRNA, complete cds. /FEA=mRNA /PROD=2-oxoglutarate carrier protein /DB_XREF=gi:3387910 /UG=Hs.184877 solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11 /FL=gb:AF070548.1	AF070548		NP_003553
209004_s_at	0.005852	gb:AF142481.1 /DEF=Homo sapiens F-box protein FLR1 (FLR1) mRNA, complete cds. /FEA=mRNA /GEN=FLR1 /PROD=F-box protein FLR1 /DB_XREF=gi:7672733 /UG=Hs.5548 f-box and leucine-rich repeat protein 5 /FL=gb:AF199420.1 gb:AF142481.1 gb:AF157323.1	AF142481		NP_277077
209005_at	0.00228	gb:AF157323.1 /DEF=Homo sapiens p45SKP2-like protein mRNA, complete cds. /FEA=mRNA /PROD=p45SKP2-like protein /DB_XREF=gi:7688696 /UG=Hs.5548 f-box and leucine-rich repeat protein 5 /FL=gb:AF199420.1 gb:AF142481.1 gb:AF157323.1	AF157323		NP_277077

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209015_s_at	1.26E-04	gb:BC002446.1 /DEF=Homo sapiens, MRJ gene for a member of the DNAJ protein family, clone MGC:1152, mRNA, complete cds. /FEA=mRNA /PROD=MRJ gene for a member of the DNAJ proteinfamily /DB_XREF=gi:12803262 /UG=Hs.181195 DnaJ (Hsp40) homolog, subfamily B, member 6 /FL=gb:BC002446.1 gb:AB015798.1	BC002446		NP_490647
209018_s_at	7.58E-04	PTEN induced putative kinase 1	BF432478	Hs.6163	NP_115785
209020_at	5.03E-04	gb:AF217514.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM038 mRNA, complete cds. /FEA=mRNA /PROD=uncharacterized bone marrow protein BM038 /DB_XREF=gi:7688970 /UG=Hs.75798 hypothetical protein /FL=gb:AF161517.1 gb:AF151041.1 gb:AF217514.1 gb:NM_016470.1	AF217514		NP_057554
209022_at	0.003171	Consensus includes gb:AK026678.1 /DEF=Homo sapiens cDNA: FLJ23025 fis, clone LNG01702, highly similar to HUMAUTOND Human autonomously replicating sequence (ARS) mRNA. /FEA=mRNA /DB_XREF=gi:10439584 /UG=Hs.8217 stromal antigen 2 /FL=gb:BC001765.1	AK026678		NP_006594
209024_s_at	0.002418	NS1-associated protein 1	AI472757	Hs.373499	NP_006363
209025_s_at	0.003171	gb:AF037448.1 /DEF=Homo sapiens RRM RNA binding protein Gry-rbp (GRY-RBP) mRNA, complete cds. /FEA=mRNA /GEN=GRY-RBP /PROD=Gry-rbp /DB_XREF=gi:3037012 /UG=Hs.155489 NS1-associated protein 1 /FL=gb:AF037448.1	AF037448		NP_006363
209027_s_at	3.26E-04	spectrin SH3 domain binding protein 1	BF673013	Hs.24752	NP_005461

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209028_s_at	0.026891	gb:AF006516.1 /DEF=Homo sapiens eps8 binding protein e3B1 mRNA, complete cds. /FEA=mRNA /PROD=e3B1 /DB_XREF=gi:2245670 /UG=Hs.24752 spectrin SH3 domain binding protein 1 /FL=gb:AF006516.1	AF006516		NP_005461
209029_at	0.003171	gb:AF193844.1 /DEF=Homo sapiens COP9 complex subunit 7a mRNA, complete cds. /FEA=mRNA /PROD=COP9 complex subunit 7a /DB_XREF=gi:6120134 /UG=Hs.3758 COP9 complex subunit 7a /FL=gb:AB033603.1 gb:AF193844.1 gb:AF210052.1 gb:NM_016319.1	AF193844		NP_057403
209033_s_at	0.001116	gb:D86550.1 /DEF=Human mRNA for serinethreonine protein kinase, complete cds. /FEA=mRNA /GEN=hMNB /PROD=serinethreonine protein kinase /DB_XREF=gi:1772437 /UG=Hs.75842 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1A /FL=gb:D85759.1 gb:D86550.1	D86550		NP_569122
209034_at	7.58E-04	gb:AF279899.1 /DEF=Homo sapiens PNAS-145 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-145 /DB_XREF=gi:12751123 /UG=Hs.75969 proline-rich protein with nuclear targeting signal /FL=gb:AF279899.1 gb:U03105.1 gb:NM_006813.1	AF279899		NP_006804

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209043_at	0.004341	gb:AF033026.1 /DEF=Homo sapiens bifunctional ATP sulfurylaseadenosine 5-phosphosulfate kinase mRNA, complete cds. /FEA=mRNA /PROD=bifunctional ATP sulfurylaseadenosine5-phosphosulfate kinase /DB_XREF=gi:3378100 /UG=Hs.3833 3-phosphoadenosine 5-phosphosulfate synthase 1 /FL=gb:AF033026.1 gb:AF016496.1 gb:NM_005443.1 gb:AF105227.1	AF033026		NP_005434
209044_x_at	0.003171	gb:BC004273.1 /DEF=Homo sapiens, splicing factor 3b, subunit 4, 49kD, clone MGC:10828, mRNA, complete cds. /FEA=mRNA /PROD=splicing factor 3b, subunit 4, 49kD /DB_XREF=gi:13279088 /UG=Hs.25797 splicing factor 3b, subunit 4, 49kD /FL=gb:BC004273.1 gb:NM_005850.1	BC004273		NP_005841
209046_s_at	1.26E-04	gb:AB030710.1 /DEF=Homo sapiens FLC3A mRNA for MAP1 light chain 3 related protein, complete cds. /FEA=mRNA /GEN=FLC3A /PROD=MAP1 light chain 3 related protein /DB_XREF=gi:12641848 /UG=Hs.6518 ganglioside expression factor 2 /FL=gb:AB030710.1 gb:AF087848.1 gb:AF077046.1 gb:NM_007285.1	AB030710		NP_009216
209050_s_at	0.026891	ral guanine nucleotide dissociation stimulator	AI421559	Hs.396157	NP_006257

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209054_s_at	0.004341	gb:AF083389.1 /DEF=Homo sapiens putative WHSC1 protein (WHSC1) mRNA, alternative splice product with complete exon 12, complete cds. /FEA=mRNA /GEN=WHSC1 /PROD=putative WHSC1 protein /DB_XREF=gi:4378016 /UG=Hs.110457 Wolf-Hirschhorn syndrome candidate 1 /FL=gb:AF083389.1	AF083389		NP_579891
209056_s_at	0.016934	CDC5 cell division cycle 5-like (S. pombe)	AW268817	Hs.155174	NP_001244
209059_s_at	0.033381	gb:AB002282.1 /DEF=Homo sapiens mRNA for hMBF1alpha, complete cds. /FEA=mRNA /PROD=hMBF1alpha /DB_XREF=gi:6526354 /UG=Hs.174050 endothelial differentiation-related factor 1 /FL=gb:NM_003792.1 gb:AB002282.1	AB002282		NP_694880
209060_x_at	0.001054	Consensus includes gb:AI438999 /FEA=EST /DB_XREF=gi:4301251 /DB_XREF=est:tc84b12.x1 /CLONE=IMAGE:2072831 /UG=Hs.225977 nuclear receptor coactivator 3 /FL=gb:AF010227.1 gb:AF012108.1 gb:AF016031.1	AF012108		NP_006525
209061_at	0.013223	Consensus includes gb:AI761748 /FEA=EST /DB_XREF=gi:5177504 /DB_XREF=est:wg67h01.x1 /CLONE=IMAGE:2370193 /UG=Hs.225977 nuclear receptor coactivator 3 /FL=gb:AF010227.1 gb:AF012108.1 gb:AF016031.1	AF012108		NP_006525
209063_x_at	0.00228	ESTs, Highly similar to polyadenylate binding protein-interacting protein 1 [Homo sapiens] [H.sapiens]	BF248165	Hs.374614	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209067_s_at	0.016934	gb:D89092.1 /DEF=Homo sapiens hnRNP JKTBP mRNA, complete cds. /FEA=mRNA /GEN=hnRNP JKTBP /DB_XREF=gi:2780747 /UG=Hs.170311 heterogeneous nuclear ribonucleoprotein D-like /FL=gb:D89092.1 gb:D89678.1	D89092		NP_112740
209073_s_at	3.26E-04	gb:AF015040.1 /DEF=Homo sapiens NUMB protein (NUMB) mRNA, complete cds. /FEA=mRNA /GEN=NUMB /PROD=NUMB protein /DB_XREF=gi:4102704 /UG=Hs.78890 numb (Drosophila) homolog /FL=gb:AF015040.1 gb:AF108092.1 gb:AF171939.1	AF015040		NP_003735
209076_s_at	0.001116	gb:BC000974.2 /DEF=Homo sapiens, Similar to hypothetical protein 628, clone MGC:5116, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein 628 /DB_XREF=gi:12803025 /UG=Hs.181349 hypothetical protein 628 /FL=gb:BC000974.2	BC000974		NP_062559
209080_x_at	0.02145	gb:AF118652.1 /DEF=Homo sapiens PKCq-interacting protein PICOT (PICOT) mRNA, complete cds. /FEA=mRNA /GEN=PICOT /PROD=PKCq-interacting protein PICOT /DB_XREF=gi:6840952 /UG=Hs.42644 thioredoxin-like /FL=gb:BC005289.1 gb:AF118649.1 gb:AF118652.1	AF118652		NP_006532
209088_s_at	6.68E-04	ubinnuclein 1	T70262	Hs.21479	NP_058632
209089_at	7.58E-04	gb:BC001267.1 /DEF=Homo sapiens, RAB5A, member RAS oncogene family, clone MGC:5048, mRNA, complete cds. /FEA=mRNA /PROD=RAB5A, member RAS oncogene family /DB_XREF=gi:12654846 /UG=Hs.73957 RAB5A, member RAS oncogene family /FL=gb:BC001267.1	BC001267		NP_004153

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209090_s_at	2.05E-04	Consensus includes gb:AL049597 /DEF=Human DNA sequence from clone RP4-612B15 on chromosome 1p22.2-31.1. Contains the (possibly pseudo) gene for a novel protein similar to 60S ribosomal protein L17 (RPL17), the gene for CGI-61, endophilin B1 and KIAA0491, ESTs, STSs, GSSs and two CpG... /FEA=mRNA_3 /DB_XREF=gi:9663352 /UG=Hs.136309 SH3-containing protein SH3GLB1 /FL=gb:AF263293.1	AL049597		
209091_s_at	1.26E-04	gb:AF263293.1 /DEF=Homo sapiens endophilin B1 mRNA, complete cds. /FEA=mRNA /PROD=endophilin B1 /DB_XREF=gi:8118529 /UG=Hs.136309 SH3-containing protein SH3GLB1 /FL=gb:AF263293.1	AF263293		NP_057093
209093_s_at	0.026891	gb:K02920.1 /DEF=Human lysosomal glucocerebrosidase mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:183021 /UG=Hs.282997 glucosidase, beta; acid (includes glucosylceramidase) /FL=gb:BC003356.1 gb:M16328.1 gb:K02920.1 gb:M19285.1 gb:NM_000157.1 gb:D13286.1	K02920		NP_000148
209096_at	0.02145	gb:U62136.2 /DEF=Homo sapiens enterocyte differentiation associated factor EDAF-1 mRNA, complete cds. /FEA=mRNA /PROD=enterocyte differentiation associated factor EDAF-1 /DB_XREF=gi:4775663 /UG=Hs.79300 ubiquitin-conjugating enzyme E2 variant 2 /FL=gb:NM_003350.2 gb:AF049140.1 gb:U62136.2	U62136		NP_003341

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209098_s_at	0.041795	gb:U61276.1 /DEF=Human transmembrane protein Jagged 1 (HJ1) mRNA, complete cds. /FEA=mRNA /GEN=HJ1 /PROD=transmembrane protein Jagged 1 /DB_XREF=gi:1438936 /UG=Hs.91143 jagged 1 (Alagille syndrome) /FL=gb:U61276.1 gb:U73936.1 gb:AF003837.1 gb:AF028593.1 gb:NM_000214.1	U61276		NP_000205
209099_x_at	1.28E-04	gb:U73936.1 /DEF=Homo sapiens Jagged 1 (HJ1) mRNA, complete cds. /FEA=mRNA /GEN=HJ1 /PROD=Jagged 1 /DB_XREF=gi:1695273 /UG=Hs.91143 jagged 1 (Alagille syndrome) /FL=gb:U61276.1 gb:U73936.1 gb:AF003837.1 gb:AF028593.1 gb:NM_000214.1	U73936		NP_000205
209102_s_at	1.26E-04	gb:AF019214.1 /DEF=Homo sapiens HMG box containing protein 1 mRNA, complete cds. /FEA=mRNA /PROD=HMG box containing protein 1 /DB_XREF=gi:2460168 /UG=Hs.10882 HMG-box containing protein 1 /FL=gb:AF019214.1	AF019214		NP_036389
209103_s_at	0.001116	gb:BC001049.1 /DEF=Homo sapiens, Similar to ubiquitin fusion degradation 1 like, clone MGC:1385, mRNA, complete cds. /FEA=mRNA /PROD=Similar to ubiquitin fusion degradation 1 like /DB_XREF=gi:12654446 /UG=Hs.181369 ubiquitin fusion degradation 1-like /FL=gb:BC001049.1 gb:BC005087.1 gb:AF141201.1	BC001049		NP_005650
209105_at	0.026891	Consensus includes gb:AI672428 /FEA=EST /DB_XREF=gi:4852159 /DB_XREF=est:wa03a11.x1 /CLONE=IMAGE:2296988 /UG=Hs.74002 nuclear receptor coactivator 1 /FL=gb:U19179.1	U19179		NP_671766

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209111_at	0.044231	gb:BC004155.1 /DEF=Homo sapiens, ring finger protein 5, clone MGC:2407, mRNA, complete cds. /FEA=mRNA /PROD=ring finger protein 5 /DB_XREF=gi:13278761 /UG=Hs.216354 ring finger protein 5 /FL=gb:BC004155.1 gb:AB056869.1 gb:NM_006913.1	BC004155		NP_008844
209112_at	0.010205	gb:BC001971.1 /DEF=Homo sapiens, Similar to cyclin-dependent kinase inhibitor 1B (p27, Kip1), clone MGC:5304, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cyclin-dependent kinase inhibitor 1B(p27, Kip1) /DB_XREF=gi:12805034 /UG=Hs.238990 cyclin-dependent kinase inhibitor 1B (p27, Kip1) /FL=gb:BC001971.1 gb:NM_004064.1 gb:U10906.1 gb:AF247551.1 gb:AY004255.1	BC001971		NP_004055
209115_at	0.010205	gb:AL117566.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566J164 (from clone DKFZp566J164); complete cds. /FEA=mRNA /GEN=DKFZp566J164 /PROD=hypothetical protein /DB_XREF=gi:5912116 /UG=Hs.154320 ubiquitin-activating enzyme E1C (homologous to yeast UBA3) /FL=gb:AF046024.1 gb:AB012190.1 gb:NM_003968.1 gb:AL117566.1	AL117566		NP_003959
209116_x_at	3.56E-04	gb:M25079.1 /DEF=Human sickle cell beta-globin mRNA, complete cds. /FEA=mRNA /PROD=beta-globin /DB_XREF=gi:179408 /UG=Hs.155376 hemoglobin, beta /FL=gb:NM_000518.3 gb:M25079.1 gb:AF117710.1 gb:AF181989.1	M25079		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209117_at	7.58E-04	gb:U79458.1 /DEF=Human WW domain binding protein-2 mRNA, complete cds. /FEA=mRNA /PROD=WW domain binding protein-2 /DB_XREF=gi:4205085 /UG=Hs.231840 WW domain binding protein 2 /FL=gb:U79458.1	U79458		
209124_at	2.05E-04	gb:U70451.1 /DEF=Human myleoid differentiation primary response protein MyD88 mRNA, complete cds. /FEA=mRNA /PROD=MyD88 /DB_XREF=gi:1763090 /UG=Hs.82116 myeloid differentiation primary response gene (88) /FL=gb:U70451.1 gb:U84408.1 gb:NM_002468.1	U70451		NP_002459
209127_s_at	0.035763	Consensus includes gb:AW173076 /FEA=EST /DB_XREF=gi:6439024 /DB_XREF=est:xj82h07.x1 /CLONE=IMAGE:2663773 /UG=Hs.116875 KIAA0156 gene product /FL=gb:AB020880.1 gb:NM_014706.1 gb:D63879.1	NM_014706		NP_055521
209129_at	0.00423	gb:AF000974.1 /DEF=Human zyxin related protein ZRP-1 mRNA, complete cds. /FEA=mRNA /PROD=ZRP-1 /DB_XREF=gi:2232135 /UG=Hs.119498 thyroid hormone receptor interactor 6 /FL=gb:BC002680.1 gb:BC004249.1 gb:BC004999.1 gb:AF000974.1	AF000974		NP_003293
209130_at	0.00161	gb:BC003686.1 /DEF=Homo sapiens, synaptosomal-associated protein, 23kD, clone MGC:5155, mRNA, complete cds. /FEA=mRNA /PROD=synaptosomal-associated protein, 23kD /DB_XREF=gi:13277555 /UG=Hs.184376 synaptosomal-associated protein, 23kD /FL=gb:BC000148.2 gb:BC003686.1 gb:U55936.1 gb:Y09567.1	BC003686		NP_570710

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209140_x_at	0.004341	gb:L42024.1 /DEF=Homo sapiens MHC HLA-B39 mRNA, complete cds. /FEA=mRNA /GEN=HLA-B39 /PROD=major histocompatibility complex /DB_XREF=gi:804748 /UG=Hs.77961 major histocompatibility complex, class I, B /FL=gb:L37880.1 gb:U29057.1 gb:U29480.1 gb:U63653.1 gb:D85761.1 gb:D85762.1 gb:M77774.1 gb:M77778.1 gb:M77777.1 gb:M16102.1 gb:M84380.1 gb:M32317.1 gb:M24033.1 gb:L17005.1 gb:U88407.1 gb:U29083.1 gb:L24373.1 gb:U03698.1 gb:U04787.1 gb:U04244.1 gb:U04245.1 gb:NM_005514.1 gb:L33922.1 gb:AF189017.1 gb:U21052.1 gb:U21053.1 gb:L36318.1 gb:U09912.1 gb:L42024.1	L42024		NP_005505
209141_at	2.05E-04	ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog, C. elegans)	AW299555	Hs.78563	NP_003333
209142_s_at	1.26E-04	gb:BC002775.1 /DEF=Homo sapiens, Similar to CG9602 gene product, clone MGC:3733, mRNA, complete cds. /FEA=mRNA /PROD=Similar to CG9602 gene product /DB_XREF=gi:12803862 /UG=Hs.78563 ubiquitin-conjugating enzyme E2G 1 (homologous to C. elegans UBC7) /FL=gb:BC002775.1 gb:NM_003342.1 gb:D78514.1	BC002775		NP_003333
209146_at	0.016934	sterol-C4-methyl oxidase-like	AV704962	Hs.239926	NP_006736
209149_s_at	0.029453	transmembrane 9 superfamily member 1	BE899402	Hs.91586	NP_006396
209150_s_at	1.26E-04	gb:U94831.1 /DEF=Homo sapiens multispanning membrane protein mRNA, complete cds. /FEA=mRNA /PROD=multispanning membrane protein /DB_XREF=gi:2276459 /UG=Hs.91586 transmembrane 9 superfamily member 1 /FL=gb:U94831.1 gb:NM_006405.1	U94831		NP_006396

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209154_at	0.007779	gb:AF234997.1 /DEF=Homo sapiens glutaminase-interacting protein 3 mRNA, complete cds. /FEA=mRNA /PROD=glutaminase-interacting protein 3 /DB_XREF=gi:12005281 /UG=Hs.12956 Tax interaction protein 1 /FL=gb:AF028823.2 gb:NM_014604.1 gb:AF234997.1	AF234997		NP_055419
209155_s_at	1.26E-04	gb:BC001595.1 /DEF=Homo sapiens, 5-nucleotidase (purine), cytosolic type B, clone MGC:1109, mRNA, complete cds. /FEA=mRNA /PROD=5-nucleotidase (purine), cytosolic type B /DB_XREF=gi:12804388 /UG=Hs.138593 5-nucleotidase (purine), cytosolic type B /FL=gb:BC001595.1	BC001595		NP_036361
209165_at	0.010205	gb:AF083208.1 /DEF=Homo sapiens Che-1 mRNA, complete cds. /FEA=mRNA /PROD=Che-1 /DB_XREF=gi:5813798 /UG=Hs.16178 apoptosis antagonizing transcription factor /FL=gb:BC000591.1 gb:AF083208.1 gb:NM_012138.1	AF083208		NP_036270
209166_s_at	0.013223	gb:U68567.1 /DEF=Human lysosomal acid alpha-mannosidase mRNA, complete cds. /FEA=mRNA /PROD=lysosomal acid alpha-mannosidase /DB_XREF=gi:1658373 /UG=Hs.279854 mannosidase, alpha, class 2B, member 1 /FL=gb:NM_000528.1 gb:BC000736.1 gb:U05572.1 gb:U68567.1 gb:U60266.1	U68567		NP_000519

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209175_at	0.013223	Consensus includes gb:AK001135.1 /DEF=Homo sapiens cDNA FLJ10273 fis, clone HEMBB1001137, highly similar to Homo sapiens mRNA for putative phospholipase. /FEA=mRNA /DB_XREF=gi:7022202 /UG=Hs.300208 Sec23-interacting protein p125 /FL=gb:BC002540.1 gb:AB019435.1 gb:NM_007190.1	AK001135		NP_009121
209185_s_at	7.58E-04	gb:AF073310.1 /DEF=Homo sapiens insulin receptor substrate-2 (IRS2) mRNA, complete cds. /FEA=mRNA /GEN=IRS2 /PROD=insulin receptor substrate-2 /DB_XREF=gi:4511968 /UG=Hs.143648 insulin receptor substrate 2 /FL=gb:NM_003749.1 gb:AF073310.1	AF073310		NP_003740
209186_at	0.00161	gb:M23114.1 /DEF=Homo sapiens calcium-ATPase (HK1) mRNA, complete cds. /FEA=mRNA /GEN=HK1 /DB_XREF=gi:184100 /UG=Hs.1526 ATPase, Ca++ transporting, cardiac muscle, slow twitch 2 /FL=gb:M23114.1	M23114		NP_733765
209187_at	7.58E-04	down-regulator of transcription 1, TBP-binding (negative cofactor 2)	AW516932	Hs.283643	NP_001929
209188_x_at	1.26E-04	gb:BC002809.1 /DEF=Homo sapiens, down-regulator of transcription 1, TBP-binding (negative cofactor 2), clone MGC:4292, mRNA, complete cds. /FEA=mRNA /PROD=down-regulator of transcription 1, TBP-binding(negative cofactor 2) /DB_XREF=gi:12803924 /UG=Hs.16697 down-regulator of transcription 1, TBP-binding (negative cofactor 2) /FL=gb:BC002809.1	BC002809		NP_001929

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209190_s_at	5.03E-04	gb:AF051782.1 /DEF=Homo sapiens diaphanous 1 (HDIA1) mRNA, complete cds. /FEA=mRNA /GEN=HDIA1 /PROD=diaphanous 1 /DB_XREF=gi:2947237 /UG=Hs.26584 diaphanous (Drosophila, homolog) 1 /FL=gb:AF051782.1 gb:NM_005219.1	AF051782		NP_005210
209194_at	3.26E-04	gb:BC005334.1 /DEF=Homo sapiens, centrin, EF-hand protein, 2, clone MGC:12421, mRNA, complete cds. /FEA=mRNA /PROD=centrin, EF-hand protein, 2 /DB_XREF=gi:13529121 /UG=Hs.82794 centrin, EF-hand protein, 2 /FL=gb:BC005334.1 gb:NM_004344.1	BC005334		NP_004335
209198_s_at	0.033381	gb:BC004291.1 /DEF=Homo sapiens, Similar to synaptotagmin 11, clone MGC:10881, mRNA, complete cds. /FEA=mRNA /PROD=Similar to synaptotagmin 11 /DB_XREF=gi:13279139 /UG=Hs.74554 KIAA0080 protein /FL=gb:BC004291.1	BC004291		NP_689493
209199_s_at	0.007779	MADS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	N22468	Hs.78995	NP_002388
209202_s_at	0.002536	gb:AF001690.1 /DEF=Homo sapiens EXT like protein 3 (EXTL3) mRNA, complete cds. /FEA=mRNA /GEN=EXTL3 /PROD=EXT like protein 3 /DB_XREF=gi:2897904 /UG=Hs.9018 exostoses (multiple)-like 3 /FL=gb:AB007042.1 gb:AF001690.1 gb:AB011091.1 gb:AF029231.1 gb:NM_001440.1 gb:AF083551.1	AF001690		NP_001431
209206_at	2.05E-04	SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)	AV701283	Hs.50785	NP_004883

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209211_at	1.49E-04	gb:AF132818.1 /DEF=Homo sapiens colon Kruppel-like factor (CKLF) mRNA, complete cds. /FEA=mRNA /GEN=CKLF /PROD=colon Kruppel-like factor /DB_XREF=gi:6580834 /UG=Hs.84728 Kruppel-like factor 5 (intestinal) /FL=gb:D14520.1 gb:NM_001730.1 gb:AF132818.1 gb:AB030824.1 gb:AF287272.1	AF132818		NP_001721
209214_s_at	0.02145	gb:BC004817.1 /DEF=Homo sapiens, Ewing sarcoma breakpoint region 1, clone MGC:5432, mRNA, complete cds. /FEA=mRNA /PROD=Ewing sarcoma breakpoint region 1 /DB_XREF=gi:13435962 /UG=Hs.129953 Ewing sarcoma breakpoint region 1 /FL=gb:BC004817.1 gb:NM_005243.1	BC004817		NP_053733
209215_at	0.005039	gb:L11669.1 /DEF=Human tetracycline transporter-like protein mRNA, complete cds. /FEA=mRNA /PROD=tetracycline transporter-like protein /DB_XREF=gi:307501 /UG=Hs.157145 tetracycline transporter-like protein /FL=gb:L11669.1 gb:NM_001120.1	L11669		NP_001111
209216_at	2.05E-04	gb:BC000464.1 /DEF=Homo sapiens, JM5 protein, clone MGC:8542, mRNA, complete cds. /FEA=mRNA /PROD=JM5 protein /DB_XREF=gi:12653388 /UG=Hs.21753 JM5 protein /FL=gb:BC000464.1 gb:BC003037.1 gb:AJ005897.1 gb:NM_007075.1	BC000464		NP_009006

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209217_s_at	1.26E-04	gb:BC000464.1 /DEF=Homo sapiens, JM5 protein, clone MGC:8542, mRNA, complete cds. /FEA=mRNA /PROD=JM5 protein /DB_XREF=gi:12653388 /UG=Hs.21753 JM5 protein /FL=gb:BC000464.1 gb:BC003037.1 gb:AJ005897.1 gb:NM_007075.1	BC000464		NP_009006
209218_at	0.00228	gb:AF098865.1 /DEF=Homo sapiens squalene epoxidase (ERG1) mRNA, complete cds. /FEA=mRNA /GEN=ERG1 /PROD=squalene epoxidase /DB_XREF=gi:4204674 /UG=Hs.71465 squalene epoxidase /FL=gb:D78130.1 gb:AF098865.1 gb:NM_003129.2	AF098865		NP_003120
209219_at	0.021312	gb:L03411.1 /DEF=Human RD protein (RD) mRNA, complete cds. /FEA=mRNA /GEN=RD /PROD=RD protein /DB_XREF=gi:190973 /UG=Hs.106061 RD RNA-binding protein /FL=gb:L03411.1 gb:NM_002904.1	L03411		NP_002895
209221_s_at	0.001116	oxysterol binding protein-like 2	AI753638	Hs.15519	NP_653081
209224_s_at	1.26E-04	gb:BC003674.1 /DEF=Homo sapiens, NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kD, B8), clone MGC:12315, mRNA, complete cds. /FEA=mRNA /PROD=NADH dehydrogenase (ubiquinone) 1 alphasubcomplex, 2 (8kD, B8) /DB_XREF=gi:13277539 /UG=Hs.163867 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kD, B8) /FL=gb:BC003674.1 gb:AF047185.1 gb:NM_002488.1 gb:AF077029.1	BC003674		NP_002479
209234_at	1.26E-04	kinesin family member 1B	BF939474	Hs.375193	NP_055889
209238_at	2.05E-04	syntaxin 3A	BE966922	Hs.82240	NP_004168

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209239_at	5.03E-04	gb:M55643.1 /DEF=Human factor KBF1 mRNA, complete cds. /FEA=mRNA /GEN=NF-kappa-B /PROD=factor KBF1 /DB_XREF=gi:189179 /UG=Hs.83428 nuclear factor of kappa light polypeptide gene enhancer in B-cells 1 (p105) /FL=gb:NM_003998.1 gb:M58603.1 gb:M55643.1	M55643		NP_003989
209247_s_at	0.028552	gb:BC001661.1 /DEF=Homo sapiens, ATP-binding cassette, sub-family F (GCN20), member 2, clone MGC:1425, mRNA, complete cds. /FEA=mRNA /PROD=ATP-binding cassette, sub-family F (GCN20),member 2 /DB_XREF=gi:12804500 /UG=Hs.153612 ATP-binding cassette, sub-family F (GCN20), member 2 /FL=gb:AF261091.1 gb:AF261092.1 gb:BC001661.1 gb:AF091073.1	BC001661		NP_009120
209248_at	0.00228	gb:AL136713.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566K113 (from clone DKFZp566K113); complete cds. /FEA=mRNA /GEN=DKFZp566K113 /PROD=hypothetical protein /DB_XREF=gi:12052945 /UG=Hs.182470 PTD010 protein /FL=gb:AF060923.1 gb:AL136713.1 gb:AF131820.1 gb:AF078863.1 gb:AB009685.1 gb:NM_014394.1	AL136713		NP_055209
209249_s_at	2.05E-04	gb:AF131820.1 /DEF=Homo sapiens clone 25077 mRNA sequence, complete cds. /FEA=mRNA /PROD=Unknown /DB_XREF=gi:4406655 /UG=Hs.182470 PTD010 protein /FL=gb:AF060923.1 gb:AL136713.1 gb:AF131820.1 gb:AF078863.1 gb:AB009685.1 gb:NM_014394.1	AF131820		NP_055209

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209250_at	0.001116	gb:BC000961.2 /DEF=Homo sapiens, degenerative spermatocyte (homolog Drosophila; lipid desaturase), clone MGC:5079, mRNA, complete cds. /FEA=mRNA /PROD=degenerative spermatocyte (homolog Drosophila; lipid desaturase) /DB_XREF=gi:12803018 /UG=Hs.185973 degenerative spermatocyte (homolog Drosophila; lipid desaturase) /FL=gb:BC000961.2	BC000961		NP_659004
209251_x_at	3.26E-04	gb:BC004949.1 /DEF=Homo sapiens, tubulin alpha 6, clone MGC:10851, mRNA, complete cds. /FEA=mRNA /PROD=tubulin alpha 6 /DB_XREF=gi:13436316 /UG=Hs.272897 Tubulin, alpha, brain-specific /FL=gb:BC004949.1	BC004949		NP_116093
209258_s_at	0.009689	Consensus includes gb:A1373676 /FEA=EST /DB_XREF=gi:4153542 /DB_XREF=est:qz53h11.x1 /CLONE=IMAGE:2030661 /UG=Hs.24485 chondroitin sulfate proteoglycan 6 (bamacan) /FL=gb:AF020043.1 gb:NM_005445.1 gb:AF067163.1	NM_005445		NP_005436
209259_s_at	0.02145	gb:AF020043.1 /DEF=Homo sapiens chromosome-associated polypeptide (HCAP) mRNA, complete cds. /FEA=mRNA /GEN=HCAP /PROD=chromosome associated polypeptide /DB_XREF=gi:3089367 /UG=Hs.24485 chondroitin sulfate proteoglycan 6 (bamacan) /FL=gb:AF020043.1 gb:NM_005445.1 gb:AF067163.1	AF020043		NP_005436
209269_s_at	6.68E-04	spleen tyrosine kinase	AW450910	Hs.74101	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209271_at	0.013223	gb:AB032251.1 /DEF=Homo sapiens BPTF mRNA for bromodomain PHD finger transcription factor, complete cds. /FEA=mRNA /GEN=BPTF /PROD=bromodomain PHD finger transcription factor /DB_XREF=gi:6683491 /UG=Hs.99872 fetal Alzheimer antigen /FL=gb:AB032251.1	AB032251		NP_004450
209273_s_at	1.41E-04	hypothetical protein MGC4276 similar to CG8198	BG387555	Hs.177776	NP_112202
209274_s_at	0.026891	gb:BC002675.1 /DEF=Homo sapiens, Similar to CG8198 gene product, clone MGC:4276, mRNA, complete cds. /FEA=mRNA /PROD=Similar to CG8198 gene product /DB_XREF=gi:12803678 /UG=Hs.177776 hypothetical protein MGC4276 similar to CG8198 /FL=gb:AF284752.1 gb:BC002675.1	BC002675		NP_112202
209276_s_at	3.26E-04	gb:AF162769.1 /DEF=Homo sapiens thioltransferase mRNA, complete cds. /FEA=mRNA /PROD=thioltransferase /DB_XREF=gi:5442445 /UG=Hs.28988 glutaredoxin (thioltransferase) /FL=gb:BC005304.1 gb:AF162769.1 gb:D21238.1	AF162769		NP_002055
209280_at	0.002864	gb:U58856.1 /DEF=Human chromosome 17 unknown product mRNA, complete cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:2209262 /UG=Hs.7835 endocytic receptor (macrophage mannose receptor family) /FL=gb:U58856.1 gb:AB014609.1 gb:AF134838.1 gb:NM_006039.1 gb:AF107292.1	U58856		NP_006030

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209281_s_at	5.03E-04	gb:M95541.1 /DEF=Homo sapiens adenosine triphosphatase mRNA, complete cds. /FEA=mRNA /PROD=adenosine triphosphatase /DB_XREF=gi:184269 /UG=Hs.78546 ATPase, Ca++ transporting, plasma membrane 1 /FL=gb:M95541.1 gb:NM_001682.1 gb:J04027.1	M95541		NP_001673
209284_s_at	1.26E-04	Consensus includes gb:AI922509 /FEA=EST /DB_XREF=gi:5658473 /DB_XREF=est:wh13g11.x1 /CLONE=IMAGE:2380676 /UG=Hs.23440 KIAA1105 protein /FL=gb:AF180425.2	AF180425		NP_056039
209285_s_at	0.00161	Consensus includes gb:N38985 /FEA=EST /DB_XREF=gi:1162192 /DB_XREF=est:yv21f02.s1 /CLONE=IMAGE:243387 /UG=Hs.23440 KIAA1105 protein /FL=gb:AF180425.2	AF180425		NP_056039
209286_at	0.001116	CDC42 effector protein (Rho GTPase binding) 3	AI754416	Hs.260024	
209287_s_at	0.02145	gb:AF104857.1 /DEF=Homo sapiens Cdc42 effector protein 3 mRNA, complete cds. /FEA=mRNA /PROD=Cdc42 effector protein 3 /DB_XREF=gi:4324453 /UG=Hs.260024 Cdc42 effector protein 3 /FL=gb:AF094521.1 gb:AF104857.1 gb:NM_006449.1 gb:AF164118.1 gb:AL136842.1	AF104857		NP_006440
209288_s_at	1.26E-04	gb:AL136842.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434A0530 (from clone DKFZp434A0530); complete cds. /FEA=mRNA /GEN=DKFZp434A0530 /PROD=hypothetical protein /DB_XREF=gi:6807668 /UG=Hs.260024 Cdc42 effector protein 3 /FL=gb:AF094521.1 gb:AF104857.1 gb:NM_006449.1 gb:AF164118.1 gb:AL136842.1	AL136842		NP_006440

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209293_x_at	0.00884	gb:U16153.1 /DEF=Human Id-4H protein mRNA, complete cds. /FEA=mRNA /PROD=Id-4H protein /DB_XREF=gi:625095 /UG=Hs.34853 inhibitor of DNA binding 4, dominant negative helix-loop-helix protein /FL=gb:NM_001546.1 gb:U16153.1 gb:U28368.1	U16153		NP_001537
209295_at	5.03E-04	gb:AF016266.1 /DEF=Homo sapiens TRAIL receptor 2 mRNA, complete cds. /FEA=mRNA /PROD=TRAIL receptor 2 /DB_XREF=gi:2529562 /UG=Hs.51233 tumor necrosis factor receptor superfamily, member 10b /FL=gb:BC001281.1 gb:AF018658.1 gb:AF016849.1 gb:AF016266.1	AF016266		NP_671716
209296_at	0.00228	gb:AF136972.1 /DEF=Homo sapiens protein phosphatase 2C-like protein mRNA, complete cds. /FEA=mRNA /PROD=protein phosphatase 2C-like protein /DB_XREF=gi:12239323 /UG=Hs.5687 protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform /FL=gb:AF136972.1 gb:AF294792.1	AF136972		NP_808908
209300_s_at	0.005852	gb:BC002888.1 /DEF=Homo sapiens, Similar to DKFZP566B183 protein, clone MGC:10356, mRNA, complete cds. /FEA=mRNA /PROD=Similar to DKFZP566B183 protein /DB_XREF=gi:12804070 /UG=Hs.12305 DKFZP566B183 protein /FL=gb:BC002888.1	BC002888		NP_056324

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209304_x_at	3.55E-04	gb:AF087853.1 /DEF=Homo sapiens growth arrest and DNA damage inducible protein beta (GADD45B) mRNA, complete cds. /FEA=mRNA /GEN=GADD45B /PROD=growth arrest and DNA damage inducible proteinbeta /DB_XREF=gi:12061050 /UG=Hs.110571 growth arrest and DNA-damage-inducible, beta /FL=gb:AF087853.1 gb:AF078077.1	AF087853		NP_056490
209308_s_at	1.26E-04	gb:BC002461.1 /DEF=Homo sapiens, BCL2adenovirus E1B 19kD-interacting protein 2, clone MGC:1529, mRNA, complete cds. /FEA=mRNA /PROD=BCL2adenovirus E1B 19kD-interacting protein 2 /DB_XREF=gi:12803290 /UG=Hs.155596 BCL2adenovirus E1B 19kD-interacting protein 2 /FL=gb:BC002461.1	BC002461		NP_004321
209310_s_at	0.041055	gb:U25804.1 /DEF=Human Ich-2 cysteine protease mRNA, complete cds. /FEA=mRNA /PROD=Ich-2 /DB_XREF=gi:886049 /UG=Hs.74122 caspase 4, apoptosis-related cysteine protease /FL=gb:U28976.1 gb:U28977.1 gb:U28978.1 gb:NM_001225.1 gb:U25804.1 gb:U28014.1	U25804		NP_150650
209311_at	2.05E-04	gb:D87461.1 /DEF=Human mRNA for KIAA0271 gene, complete cds. /FEA=mRNA /GEN=KIAA0271 /DB_XREF=gi:1944417 /UG=Hs.75244 BCL2-like 2 /FL=gb:U59747.1 gb:D87461.1 gb:NM_004050.1	D87461		NP_004041

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209312_x_at	0.010205	gb:U65585.1 /DEF=Homo sapiens MHC class II antigen (HLA-DRB1) mRNA, HLA-DRB1*PBL allele, complete cds. /FEA=mRNA /GEN=HLA-DRB1 /PROD=MHC class II antigen /DB_XREF=gi:5478215 /UG=Hs.180255 major histocompatibility complex, class II, DR beta 1 /FL=gb:M33600.1 gb:NM_002124.1 gb:M28583.1 gb:U65585.1	U65585		NP_002115
209326_at	0.00161	gb:D84454.1 /DEF=Human mRNA for UDP-galactose translocator, complete cds. /FEA=mRNA /PROD=UDP-galactose translocator /DB_XREF=gi:1526437 /UG=Hs.21899 solute carrier family 35 (UDP-galactose transporter), member 2 /FL=gb:D84454.1	D84454		NP_005651
209333_at	6.68E-04	gb:AB018265.1 /DEF=Homo sapiens mRNA for KIAA0722 protein, complete cds. /FEA=mRNA /GEN=KIAA0722 /PROD=KIAA0722 protein /DB_XREF=gi:3882164 /UG=Hs.47061 unc-51 (C. elegans)-like kinase 1 /FL=gb:AF045458.1 gb:AB018265.1 gb:NM_003565.1	AB018265		NP_003556
209338_at	1.26E-04	gb:U03494.1 /DEF=Human transcription factor LSF mRNA, complete cds. /FEA=mRNA /PROD=transcription factor LSF /DB_XREF=gi:476098 /UG=Hs.154970 transcription factor CP2 /FL=gb:BC003634.1 gb:U03494.1	U03494		NP_005644
209339_at	3.56E-04	gb:U76248.1 /DEF=Human hSIAH2 mRNA, complete cds. /FEA=mRNA /PROD=hSIAH2 /DB_XREF=gi:2673967 /UG=Hs.20191 seven in absentia (Drosophila) homolog 2 /FL=gb:U76248.1 gb:NM_005067.1	U76248		NP_005058

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209344_at	0.041055	gb:BC002827.1 /DEF=Homo sapiens, tropomyosin 4, clone MGC:3641, mRNA, complete cds. /FEA=mRNA /PROD=tropomyosin 4 /DB_XREF=gi:12803958 /UG=Hs.250641 tropomyosin 4 /FL=gb:BC002827.1 gb:NM_003290.1	BC002827		NP_003281
209357_at	1.26E-04	gb:AF109161.1 /DEF=Homo sapiens p35srj (MRG1) mRNA, complete cds. /FEA=mRNA /GEN=MRG1 /PROD=p35srj /DB_XREF=gi:4193945 /UG=Hs.82071 Cbpp300-interacting transactivator, with GluAsp-rich carboxy-terminal domain, 2 /FL=gb:BC004377.1 gb:AF109161.1	AF109161		NP_006070
209367_at	0.00228	gb:AB002559.1 /DEF=Homo sapiens mRNA for hunc18b2, complete cds. /FEA=mRNA /PROD=hunc18b2 /DB_XREF=gi:1944129 /UG=Hs.90535 syntaxin binding protein 2 /FL=gb:BC002869.1 gb:U63533.1 gb:AB002559.1 gb:NM_006949.1	AB002559		NP_008880
209369_at	1.26E-04	gb:M63310.1 /DEF=Human 1,2-cyclic-inositol-phosphate phosphodiesterase (ANX3) mRNA, complete cds. /FEA=mRNA /GEN=ANX3 /PROD=1,2-cyclic-inositol-phosphate phosphodiesterase /DB_XREF=gi:178696 /UG=Hs.1378 annexin A3 /FL=gb:BC000871.1 gb:M63310.1 gb:M20560.1 gb:NM_005139.1	M63310		NP_005130

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209375_at	0.010205	gb:D21089.1 /DEF=Human mRNA for XP-C repair complementing protein (p125), complete cds. /FEA=mRNA /GEN=XPCC /PROD=XP-C repair complementing protein (p125) /DB_XREF=gi:475156 /UG=Hs.320 xeroderma pigmentosum, complementation group C /FL=gb:D21089.1 gb:NM_004628.1	D21089		NP_004619
209376_x_at	6.68E-04	splicing factor, arginine/serine-rich 2, interacting protein	AW084759	Hs.51957	NP_004710
209380_s_at	0.00161	gb:AF146074.1 /DEF=Homo sapiens ABC protein mRNA, complete cds. /FEA=mRNA /PROD=ABC protein /DB_XREF=gi:5006890 /UG=Hs.108660 ATP-binding cassette, sub-family C (CFTRMRP), member 5 /FL=gb:AF104942.1 gb:AB019002.1 gb:AF146074.1 gb:NM_005688.1 gb:U83661.2	AF146074		NP_005679
209382_at	0.016934	gb:U93867.1 /DEF=Human RNA polymerase III subunit (RPC62) mRNA, complete cds. /FEA=mRNA /GEN=RPC62 /PROD=RNA polymerase III subunit /DB_XREF=gi:2228747 /UG=Hs.250745 polymerase (RNA) III (DNA directed) (62kD) /FL=gb:BC002586.1 gb:NM_006468.2 gb:U93867.1	U93867		NP_006459
209384_at	5.03E-04	proline synthetase co-transcribed homolog (bacterial)	AA176833	Hs.301959	NP_009129
209388_at	0.004341	gb:BC000927.1 /DEF=Homo sapiens, Similar to poly (A) polymerase, clone MGC:5378, mRNA, complete cds. /FEA=mRNA /PROD=Similar to poly (A) polymerase /DB_XREF=gi:12654216 /UG=Hs.49007 poly(A) polymerase alpha /FL=gb:BC000927.1	BC000927		NP_116021

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209390_at	0.016934	gb:AF013168.1 /DEF=Homo sapiens hamartin (TSC1) mRNA, complete cds. /FEA=mRNA /GEN=TSC1 /PROD=hamartin /DB_XREF=gi:2331280 /UG=Hs.79393 tuberous sclerosis 1 (hamartin) /FL=gb:AF013168.1 gb:NM_000368.1	AF013168		NP_000359
209392_at	0.008237	gb:L35594.1 /DEF=Human autotaxin mRNA, complete cds. /FEA=mRNA /PROD=autotaxin /DB_XREF=gi:537905 /UG=Hs.174185 ectonucleotide pyrophosphatasephosphodiesterase 2 (autotaxin) /FL=gb:L35594.1	L35594		NP_006200
209393_s_at	1.26E-04	gb:AF047695.1 /DEF=Homo sapiens cap-binding protein 4EHP mRNA, complete cds. /FEA=mRNA /PROD=cap-binding protein 4EHP /DB_XREF=gi:3172338 /UG=Hs.19122 eukaryotic translation initiation factor 4E-like 3 /FL=gb:BC005392.1 gb:AF047695.1 gb:AF068117.1 gb:AF038957.1 gb:NM_004846.1	AF047695		NP_004837
209397_at	1.26E-04	gb:BC000147.1 /DEF=Homo sapiens, Similar to malic enzyme 2, NAD(+)-dependent, mitochondrial, clone MGC:5187, mRNA, complete cds. /FEA=mRNA /PROD=Similar to malic enzyme 2, NAD(+)-dependent, mitochondrial /DB_XREF=gi:12652790 /UG=Hs.75342 malic enzyme 2, NAD(+)-dependent, mitochondrial /FL=gb:BC000147.1	BC000147		NP_002387

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209403_at	0.026891	gb:AL136860.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434P2235 (from clone DKFZp434P2235); complete cds. /FEA=mRNA /GEN=DKFZp434P2235 /PROD=hypothetical protein /DB_XREF=gi:12053224 /UG=Hs.105891 Homo sapiens mRNA; cDNA DKFZp434P2235 (from clone DKFZp434P2235); complete cds /FL=gb:AL136860.1	AL136860		NP_115634
209404_s_at	2.05E-04	gb:AF151867.1 /DEF=Homo sapiens CGI-109 protein mRNA, complete cds. /FEA=mRNA /PROD=CGI-109 protein /DB_XREF=gi:4929686 /UG=Hs.278391 CGI-109 protein /FL=gb:AF151867.1	AF151867		
209409_at	1.26E-04	gb:D86962.1 /DEF=Human mRNA for KIAA0207 gene, complete cds. /FEA=mRNA /GEN=KIAA0207 /DB_XREF=gi:1503997 /UG=Hs.81875 growth factor receptor-bound protein 10 /FL=gb:D86962.1 gb:AF000017.1	D86962		NP_005302
209411_s_at	0.002418	golgi associated, gamma adaptin ear containing, ARF binding protein 3	AW008018	Hs.87726	NP_619525
209418_s_at	0.001116	gb:BC003615.1 /DEF=Homo sapiens, gene from NF2meningioma region of 22q12, clone MGC:1540, mRNA, complete cds. /FEA=mRNA /PROD=gene from NF2meningioma region of 22q12 /DB_XREF=gi:13177658 /UG=Hs.75361 gene from NF2meningioma region of 22q12 /FL=gb:BC003615.1 gb:NM_003678.1 gb:AB023200.1	BC003615		NP_003669

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209424_s_at	0.003817	Consensus includes gb:AI796120 /FEA=EST /DB_XREF=gi:5361583 /DB_XREF=est:wh42f03.x1 /CLONE=IMAGE:2383421 /UG=Hs.128749 alpha-methylacyl-CoA racemase /FL=gb:AF047020.1 gb:AF158378.1 gb:NM_014324.1	NM_014324		NP_055139
209426_s_at	0.01159	gb:AF047020.1 /DEF=Homo sapiens alpha-methylacyl-CoA racemase mRNA, complete cds. /FEA=mRNA /PROD=alpha-methylacyl-CoA racemase /DB_XREF=gi:4204096 /UG=Hs.128749 alpha-methylacyl-CoA racemase /FL=gb:AF047020.1 gb:AF158378.1 gb:NM_014324.1	AF047020		NP_055139
209429_x_at	0.016686	gb:AF112207.1 /DEF=Homo sapiens translation initiation factor eIF-2b delta subunit mRNA, complete cds. /FEA=mRNA /PROD=translation initiation factor eIF-2b deltasubunit /DB_XREF=gi:6563201 /UG=Hs.169474 DKFZP586J0119 protein /FL=gb:BC001870.1 gb:AF112207.1	AF112207		NP_056451
209438_at	0.004341	Consensus includes gb:AL096700 /DEF=Human DNA sequence from clone RP3-499B10 on chromosome Xp22.31-22.13. Contains the 3 end of the gene for serinethreonine protein phosphatase PPEF, the 3 end of the PHKA2 gene for liver phosphorylase kinase alpha 2, ESTs, STSs and GSSs /FEA=mRNA_2 /DB_XREF=gi:7077104 /UG=Hs.54941 phosphorylase kinase, alpha 2 (liver) /FL=gb:D38616.1 gb:NM_000292.1	AL096700		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209439_s_at	5.03E-04	gb:D38616.1 /DEF=Human mRNA for phosphorylase kinase alpha subunit, complete cds. /FEA=mRNA /PROD=phosphorylase kinase alpha subunit /DB_XREF=gi:1304117 /UG=Hs.54941 phosphorylase kinase, alpha 2 (liver) /FL=gb:D38616.1 gb:NM_000292.1	D38616		NP_000283
209447_at	0.005852	gb:AF043290.1 /DEF=Homo sapiens lymphocyte membrane associated protein (8B7) mRNA, complete cds. /FEA=mRNA /GEN=8B7 /PROD=lymphocyte membrane associated protein /DB_XREF=gi:2895592 /UG=Hs.8182 synaptic nuclei expressed gene 1b /FL=gb:AF043290.1	AF043290		NP_598411
209448_at	1.26E-04	gb:BC002439.1 /DEF=Homo sapiens, Tat-interacting protein (30kD), clone MGC:1318, mRNA, complete cds. /FEA=mRNA /PROD=Tat-interacting protein (30kD) /DB_XREF=gi:12803250 /UG=Hs.90753 Tat-interacting protein (30kD) /FL=gb:BC002439.1 gb:AF039103.1	BC002439		NP_006401
209451_at	0.013223	gb:U59863.1 /DEF=Human TRAF-interacting protein I-TRAF mRNA, complete cds. /FEA=mRNA /PROD=I-TRAF /DB_XREF=gi:1518017 /UG=Hs.146847 TRAF family member-associated NFKB activator /FL=gb:U59863.1	U59863		NP_597841

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209452_s_at	3.26E-04	gb:AF035824.1 /DEF=Homo sapiens vesicle soluble NSF attachment protein receptor (VT11) mRNA, complete cds. /FEA=mRNA /GEN=VTI1 /PROD=vesicle soluble NSF attachment protein receptor /DB_XREF=gi:2687399 /UG=Hs.169206 vesicle-associated soluble NSF attachment protein receptor (v-SNARE; homolog of S. cerevisiae VT11) /FL=gb:BC003142.1 gb:AF035824.1 gb:AF060902.1 gb:NM_006370.1	AF035824		NP_006361
209453_at	0.003166	gb:M81768.1 /DEF=Human NaH antiporter (APNH1) mRNA, complete cds. /FEA=mRNA /GEN=APNH /PROD=NaH antiporter /DB_XREF=gi:178752 /UG=Hs.170222 solute carrier family 9 (sodiumhydrogen exchanger), isoform 1 (antiporter, Na+H+, amiloride sensitive) /FL=gb:M81768.1 gb:AF141350.1 gb:AF141351.1 gb:AF141352.1 gb:AF141353.1 gb:AF141354.1 gb:AF141355.1 gb:AF141356.1 gb:AF141357.1 gb:AF141358.1 gb:AF141359.1 gb:AF146430.1 gb:AF146431.1 gb:AF146432.1 gb:AF146433.1 gb:AF146434.1 gb:AF146435.1 gb:AF146436.1 gb:AF146437.1 gb:AF146438.1 gb:AF146439.1	M81768		NP_003038
209455_at	1.26E-04	F-box and WD-40 domain protein 1B	BE963245	Hs.21229	NP_387449
209457_at	0.033381	gb:U16996.1 /DEF=Human protein tyrosine phosphatase mRNA, complete cds. /FEA=mRNA /PROD=protein tyrosine phosphatase /DB_XREF=gi:642012 /UG=Hs.2128 dual specificity phosphatase 5 /FL=gb:NM_004419.2 gb:U16996.1 gb:U15932.2	U16996		NP_004410

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209458_x_at	3.26E-04	gb:AF105974.1 /DEF=Homo sapiens alpha one globin (HBA1) mRNA, complete cds. /FEA=mRNA /GEN=HBA1 /PROD=alpha one globin /DB_XREF=gi:4038449 /UG=Hs.272572 hemoglobin, alpha 2 /FL=gb:AF097635.1 gb:AF105974.1 gb:NM_000517.2	AF105974		NP_000549
209459_s_at	0.041055	gb:AF237813.1 /DEF=Homo sapiens NPD009 mRNA, complete cds. /FEA=mRNA /PROD=NPD009 /DB_XREF=gi:9963907 /UG=Hs.283675 NPD009 protein /FL=gb:NM_020686.1 gb:AF237813.1	AF237813		NP_065737
209463_s_at	0.001713	gb:D50544.1 /DEF=Human TFIID subunit p22 mRNA, complete cds. /FEA=mRNA /PROD=TFIID subunit p22 /DB_XREF=gi:1345403 /UG=Hs.82037 TATA box binding protein (TBP)-associated factor, RNA polymerase II, J, 20kD /FL=gb:D50544.1 gb:NM_005644.2	D50544		NP_005635
209467_s_at	5.03E-04	gb:BC002755.1 /DEF=Homo sapiens, Similar to MAP kinase-interacting serinethreonine kinase 1, clone MGC:3690, mRNA, complete cds. /FEA=mRNA /PROD=Similar to MAP kinase-interactingserinethreonine kinase 1 /DB_XREF=gi:12803828 /UG=Hs.5591 MAP kinase-interacting serinethreonine kinase 1 /FL=gb:BC002755.1	BC002755		NP_003675
209471_s_at	0.00228	gb:L00634.1 /DEF=Human farnesyl-protein transferase alpha-subunit mRNA, complete cds. /FEA=mRNA /PROD=farnesyl-protein transferase alpha-subunit /DB_XREF=gi:292030 /UG=Hs.138381 farnesyltransferase, CAAX box, alpha /FL=gb:L00634.1 gb:L10413.1 gb:NM_002027.1	L00634		NP_002018

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209472_at	0.007779	gb:BC000819.1 /DEF=Homo sapiens, Similar to CG6950 gene product, clone MGC:5114, mRNA, complete cds. /FEA=mRNA /PROD=Similar to CG6950 gene product /DB_XREF=gi:12654030 /UG=Hs.180378 hypothetical protein 669 /FL=gb:BC000819.1 gb:NM_019610.1	BC000819		NP_062556
209473_at	1.26E-04	ectonucleoside triphosphate diphosphohydrolase 1	AV717590	Hs.205353	NP_001767
209475_at	0.004341	gb:AF106069.1 /DEF=Homo sapiens deubiquitinating enzyme (UNPH4) mRNA, complete cds. /FEA=mRNA /GEN=UNPH4 /PROD=deubiquitinating enzyme /DB_XREF=gi:5814096 /UG=Hs.23168 ubiquitin specific protease 15 /FL=gb:AF013990.1 gb:AF106069.1	AF106069		NP_006304
209476_at	5.03E-04	Consensus includes gb:AL080080.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564E1962 (from clone DKFZp564E1962); partial cds. /FEA=mRNA /GEN=DKFZp564E1962 /PROD=hypothetical protein /DB_XREF=gi:5262491 /UG=Hs.24766 thioredoxin-related transmembrane protein /FL=gb:AB048246.1	AL080080		NP_110382
209479_at	1.26E-04	gb:BC000758.1 /DEF=Homo sapiens, clone MGC:2698, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:2698) /DB_XREF=gi:12653928 /UG=Hs.44468 DKFZP586D0623 protein /FL=gb:BC000758.1	BC000758		NP_056254

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209480_at	0.022752	gb:M16276.1 /DEF=Human MHC class II HLA-DR2-Dw12 mRNA DQw1-beta, complete cds. /FEA=mRNA /GEN=HLA-DRB2 /DB_XREF=gi:188397 /UG=Hs.73931 major histocompatibility complex, class II, DQ beta 1 /FL=gb:M60028.1 gb:M17564.1 gb:M81140.1 gb:M81141.1 gb:M16276.1 gb:NM_002123.1	M16276		NP_002114
209481_at	0.005852	gb:AF226044.1 /DEF=Homo sapiens HSNFRK (HSNFRK) mRNA, complete cds. /FEA=mRNA /GEN=HSNFRK /PROD=HSNFRK /DB_XREF=gi:9295326 /UG=Hs.79025 KIAA0096 protein /FL=gb:AF226044.1	AF226044		NP_060189
209482_at	0.005339	gb:BC001430.1 /DEF=Homo sapiens, POP7 (processing of precursor, S. cerevisiae) homolog, clone MGC:1986, mRNA, complete cds. /FEA=mRNA /PROD=POP7 (processing of precursor, S. cerevisiae)homolog /DB_XREF=gi:12655150 /UG=Hs.18747 POP7 (processing of precursor, S. cerevisiae) homolog /FL=gb:BC001430.1 gb:U94316.1 gb:NM_005837.1	BC001430		NP_005828
209497_s_at	0.02145	gb:BC003503.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 4921506I22 gene, clone MGC:10380, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 4921506I22 gene /DB_XREF=gi:13097557 /UG=Hs.49994 Homo sapiens, clone MGC:10871, mRNA, complete cds /FL=gb:BC003503.1 gb:BC004951.1	BC003503		NP_113680

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209498_at	3.56E-04	Consensus includes gb:X16354.1 /DEF=Human mRNA for transmembrane carcinoembryonic antigen BGP _a (formerly TM1-CEA). /FEA=mRNA /PROD=TM1-CEA preprotein /DB_XREF=gi:37197 /UG=Hs.50964 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein) /FL=gb:J03858.1	X16354		NP_001703
209500_x_at	0.00228	gb:AF114012.1 /DEF=Homo sapiens tumor necrosis factor-related death ligand-1beta mRNA, complete cds. /FEA=mRNA /PROD=tumor necrosis factor-related deathligand-1beta /DB_XREF=gi:7328555 /UG=Hs.54673 tumor necrosis factor (ligand) superfamily, member 13 /FL=gb:AF114012.1	AF114012		NP_742086
209510_at	0.007779	gb:AF064801.1 /DEF=Homo sapiens multiple membrane spanning receptor TRC8 (TRC8) mRNA, complete cds. /FEA=mRNA /GEN=TRC8 /PROD=multiple membrane spanning receptor TRC8 /DB_XREF=gi:3395786 /UG=Hs.28285 patched related protein translocated in renal cancer /FL=gb:AF064801.1 gb:NM_007218.1	AF064801		NP_009149
209512_at	3.26E-04	gb:BC004331.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 2610207I16 gene, clone MGC:10940, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 2610207I16 gene /DB_XREF=gi:13279253 /UG=Hs.47986 Homo sapiens, Similar to RIKEN cDNA 2610207I16 gene, clone MGC:10940, mRNA, complete cds /FL=gb:BC004331.1	BC004331		NP_115679

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209513_s_at	3.26E-04	gb:BC004331.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 2610207I16 gene, clone MGC:10940, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 2610207I16 gene /DB_XREF=gi:13279253 /UG=Hs.47986 Homo sapiens, Similar to RIKEN cDNA 2610207I16 gene, clone MGC:10940, mRNA, complete cds /FL=gb:BC004331.1	BC004331		NP_115679
209514_s_at	0.00161	RAB27A, member RAS oncogene family	BE502030	Hs.50477	NP_004571
209515_s_at	2.05E-04	gb:U38654.3 /DEF=Homo sapiens Rab27a mRNA, complete cds. /FEA=mRNA /PROD=Rab27a /DB_XREF=gi:4887230 /UG=Hs.50477 RAB27A, member RAS oncogene family /FL=gb:U38654.3	U38654		NP_004571
209517_s_at	3.26E-04	gb:AB020982.1 /DEF=Homo sapiens ASH2L mRNA, complete cds, similar to Drosophila ash2 sequence. /FEA=mRNA /GEN=ASH2L /DB_XREF=gi:4417209 /UG=Hs.6856 ash2 (absent, small, or homeotic, Drosophila, homolog)-like /FL=gb:AF056718.1 gb:AB020982.1 gb:NM_004674.1	AB020982		NP_004665
209520_s_at	0.007779	gb:BC001450.1 /DEF=Homo sapiens, nuclear cap binding protein subunit 1, 80kD, clone MGC:2087, mRNA, complete cds. /FEA=mRNA /PROD=nuclear cap binding protein subunit 1, 80kD /DB_XREF=gi:12655186 /UG=Hs.89563 nuclear cap binding protein subunit 1, 80kD /FL=gb:BC001450.1 gb:D32002.1 gb:NM_002486.1	BC001450		NP_002477

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209525_at	0.012818	Consensus includes gb:BG285017 /FEA=EST /DB_XREF=gi:13036553 /DB_XREF=est:602409285F1 /CLONE=IMAGE:4538229 /UG=Hs.127842 CGI-142 /FL=gb:AF151900.1 gb:AB029156.1 gb:NM_016073.1	NM_016073		NP_057157
209539_at	0.013223	Consensus includes gb:D25304.1 /DEF=Human mRNA for KIAA0006 gene, partial cds. /FEA=mRNA /GEN=KIAA0006 /DB_XREF=gi:435445 /UG=Hs.79307 RacCdc42 guanine exchange factor (GEF) 6 /FL=gb:D13631.1	D25304		NP_004831
209549_s_at	7.58E-04	gb:BC001121.1 /DEF=Homo sapiens, Similar to deoxyguanosine kinase, clone MGC:2111, mRNA, complete cds. /FEA=mRNA /PROD=Similar to deoxyguanosine kinase /DB_XREF=gi:12654572 /UG=Hs.326494 Homo sapiens, Similar to deoxyguanosine kinase, clone MGC:2111, mRNA, complete cds /FL=gb:BC001121.1	BC001121		NP_550440
209551_at	3.26E-04	gb:BC004875.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 2310034L04 gene, clone MGC:11061, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 2310034L04 gene /DB_XREF=gi:13436109 /UG=Hs.66309 Homo sapiens, Similar to RIKEN cDNA 2310034L04 gene, clone MGC:11061, mRNA, complete cds /FL=gb:BC004875.1	BC004875		NP_115688

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209553_at	1.26E-04	gb:BC001001.2 /DEF=Homo sapiens, Similar to hBKLF for basic kruppel like factor, clone MGC:5362, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hBKLF for basic kruppel like factor /DB_XREF=gi:12803042 /UG=Hs.7316 KIAA0804 protein /FL=gb:BC001001.2	BC001001		
209555_s_at	0.00161	gb:M98399.1 /DEF=Human antigen CD36 (clone 21) mRNA, complete cds. /FEA=mRNA /GEN=CD36 /PROD=antigen CD36 /DB_XREF=gi:180112 /UG=Hs.75613 CD36 antigen (collagen type I receptor, thrombospondin receptor) /FL=gb:M98399.1	M98399		NP_000063
209561_at	0.020456	gb:L38969.1 /DEF=Homo sapiens thrombospondin 3 (THBS3) mRNA, complete cds. /FEA=mRNA /GEN=THBS3 /PROD=thrombospondin 3 /DB_XREF=gi:886298 /UG=Hs.169875 thrombospondin 3 /FL=gb:NM_007112.1 gb:L38969.1	L38969		NP_009043
209566_at	0.001116	Consensus includes gb:AL080184.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434O071 (from clone DKFZp434O071). /FEA=mRNA /DB_XREF=gi:5262661 /UG=Hs.7089 insulin induced protein 2 /FL=gb:AF125392.1	AL080184		NP_057217
209574_s_at	0.00161	chromosome 18 open reading frame 1	AI349506	Hs.153498	NP_004329

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209575_at	0.010205	gb:BC001903.1 /DEF=Homo sapiens, Similar to interleukin 10 receptor, beta, clone MGC:2210, mRNA, complete cds. /FEA=mRNA /PROD=Similar to interleukin 10 receptor, beta /DB_XREF=gi:12804902 /UG=Hs.173936 interleukin 10 receptor, beta /FL=gb:BC001903.1 gb:NM_000628.1	BC001903		NP_000619
209579_s_at	0.00161	methyl-CpG binding domain protein 4	AL556619	Hs.35947	NP_003916
209580_s_at	0.004341	gb:AF114784.1 /DEF=Homo sapiens methyl-CpG binding endonuclease (MED1) mRNA, complete cds. /FEA=mRNA /GEN=MED1 /PROD=methyl-CpG binding endonuclease /DB_XREF=gi:4539758 /UG=Hs.35947 methyl-CpG binding domain protein 4 /FL=gb:AF072250.1 gb:NM_003925.1 gb:AF114784.1	AF114784		NP_003916
209593_s_at	0.00228	gb:AF317129.1 /DEF=Homo sapiens FKSG18 (FKSG18) mRNA, complete cds. /FEA=mRNA /GEN=FKSG18 /PROD=FKSG18 /DB_XREF=gi:12276123 /UG=Hs.252682 torsin family 1, member B (torsin B) /FL=gb:AF317129.1	AF317129		NP_055321
209595_at	0.002313	gb:BC001771.1 /DEF=Homo sapiens, general transcription factor IIF, polypeptide 2 (30kD subunit), clone MGC:1502, mRNA, complete cds. /FEA=mRNA /PROD=general transcription factor IIF, polypeptide 2(30kD subunit) /DB_XREF=gi:12804688 /UG=Hs.58593 general transcription factor IIF, polypeptide 2 (30kD subunit) /FL=gb:BC001771.1 gb:NM_004128.1	BC001771		NP_004119
209599_s_at	0.045257	TcD37 homolog	AI762105	Hs.332683	NP_067045

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209604_s_at	0.00228	gb:BC003070.1 /DEF=Homo sapiens, GATA-binding protein 3, clone MGC:2346, mRNA, complete cds. /FEA=mRNA /PROD=GATA-binding protein 3 /DB_XREF=gi:13111765 /UG=Hs.169946 GATA-binding protein 3 /FL=gb:BC003070.1 gb:M69106.1 gb:NM_002051.1	BC003070		NP_002042
209605_at	0.007943	gb:D87292.1 /DEF=Homo sapiens mRNA for rhodanese, complete cds. /FEA=mRNA /PROD=rhodanese /DB_XREF=gi:1877030 /UG=Hs.248267 thiosulfate sulfurtransferase (rhodanese) /FL=gb:D87292.1	D87292		NP_003303
209610_s_at	5.64E-04	solute carrier family 1 (glutamate/neutral amino acid transporter), member 4	BF340083	Hs.323878	NP_003029
209615_s_at	0.004341	gb:U51120.1 /DEF=Human protein kinase PAK1 mRNA, complete cds. /FEA=mRNA /PROD=PAK1 /DB_XREF=gi:1256421 /UG=Hs.62402 p21Cdc42Rac1-activated kinase 1 (yeast Ste20-related) /FL=gb:U51120.1 gb:NM_002576.2 gb:U24152.1	U51120		NP_002567
209619_at	3.26E-04	gb:K01144.1 /DEF=Human major histocompatibility class II antigen gamma chain mRNA, complete cds. /FEA=mRNA /PROD=class II antigen gamma chain /DB_XREF=gi:188469 /UG=Hs.84298 CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated) /FL=gb:NM_004355.1 gb:K01144.1	K01144		NP_004346

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209620_s_at	0.022752	gb:AB005289.1 /DEF=Homo sapiens mRNA for ABC transporter 7 protein, complete cds. /FEA=mRNA /GEN=hABC7 /PROD=ABC transporter 7 protein /DB_XREF=gi:3228278 /UG=Hs.125856 ATP-binding cassette, sub-family B (MDRTAP), member 7 /FL=gb:AB005289.1 gb:AF038950.1 gb:AF133659.1 gb:NM_004299.2	AB005289		NP_004290
209628_at	0.007779	Consensus includes gb:AK023289.1 /DEF=Homo sapiens cDNA FLJ13227 fis, clone OVARC1000071, weakly similar to Homo sapiens NTF2-related export protein NXT1 (NXT1) mRNA. /FEA=mRNA /DB_XREF=gi:10435160 /UG=Hs.25010 hypothetical protein P15-2 /FL=gb:AF246127.1 gb:NM_018698.1 gb:AF201942.1	AK023289		NP_061168
209630_s_at	1.26E-04	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AL043967	Hs.350947	
209640_at	3.26E-04	gb:M79462.1 /DEF=Human PML-1 mRNA, complete CDS. /FEA=mRNA /GEN=PML /DB_XREF=gi:190116 /UG=Hs.89633 promyelocytic leukemia /FL=gb:M79462.1	M79462		NP_150253
209643_s_at	0.002537	gb:AF033850.1 /DEF=Homo sapiens phospholipase D2 (PLD2) mRNA, complete cds. /FEA=mRNA /GEN=PLD2 /PROD=phospholipase D2 /DB_XREF=gi:2645857 /UG=Hs.104519 phospholipase D2 /FL=gb:AF033850.1 gb:AF038440.1 gb:AF035483.1 gb:NM_002663.1	AF033850		NP_002654

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209649_at	0.005339	gb:AL133600.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564C047 (from clone DKFZp564C047); complete cds. /FEA=mRNA /GEN=DKFZp564C047 /PROD=hypothetical protein /DB_XREF=gi:6599205 /UG=Hs.17200 STAM-like protein containing SH3 and ITAM domains 2 /FL=gb:AF042273.1 gb:AL133600.1	AL133600		NP_005834
209653_at	6.93E-04	gb:U93240.1 /DEF=Human importin alpha 3 mRNA, complete cds. /FEA=mRNA /PROD=importin alpha 3 /DB_XREF=gi:1928974 /UG=Hs.302499 karyopherin alpha 4 (importin alpha 3) /FL=gb:AB002533.2 gb:NM_002268.2 gb:U93240.1	U93240		NP_002259
209665_at	0.041055	gb:AF040704.1 /DEF=Homo sapiens putative tumor suppressor protein (101F6) mRNA, complete cds. /FEA=mRNA /GEN=101F6 /PROD=putative tumor suppressor protein /DB_XREF=gi:2935319 /UG=Hs.149443 putative tumor suppressor /FL=gb:AF040704.1 gb:NM_007022.1	AF040704		NP_008953
209666_s_at	2.05E-04	gb:AF080157.1 /DEF=Homo sapiens Ikb kinase-a (IKK-alpha) mRNA, complete cds. /FEA=mRNA /GEN=IKK-alpha /PROD=Ikb kinase-a /DB_XREF=gi:4185272 /UG=Hs.198998 conserved helix-loop-helix ubiquitous kinase /FL=gb:U22512.1 gb:AF012890.1 gb:AF009225.1 gb:AF080157.1 gb:NM_001278.1	AF080157		NP_001269
209667_at	1.32E-04	carboxylesterase 2 (intestine, liver)	BF033242	Hs.282975	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209670_at	0.005852	gb:M12959.1 /DEF=Human T-cell receptor active alpha-chain mRNA from JM cell line, complete cds. /FEA=mRNA /GEN=TCRA /DB_XREF=gi:338734 /UG=Hs.74647 Human T-cell receptor active alpha-chain mRNA from JM cell line, complete cds /FL=gb:M12959.1 gb:M12423.1	M12959		
209671_x_at	0.001618	gb:M12423.1 /DEF=Human T-cell receptor active alpha-chain mRNA, complete cds, from Jurkat T-cell line. /FEA=mRNA /GEN=TCRA /DB_XREF=gi:338738 /UG=Hs.74647 Human T-cell receptor active alpha-chain mRNA from JM cell line, complete cds /FL=gb:M12959.1 gb:M12423.1	M12423		
209678_s_at	2.05E-04	gb:L18964.1 /DEF=Human protein kinase C iota isoform (PRKCI) mRNA, complete cds. /FEA=mRNA /GEN=PRKCI /PROD=protein kinase C iota /DB_XREF=gi:432273 /UG=Hs.1904 protein kinase C, iota /FL=gb:L18964.1 gb:NM_002740.1 gb:L33881.1	L18964		NP_002731
209681_at	0.004733	gb:AF153330.1 /DEF=Homo sapiens thiamine carrier 1 (TC1) mRNA, complete cds. /FEA=mRNA /GEN=TC1 /PROD=thiamine carrier 1 /DB_XREF=gi:5453325 /UG=Hs.30246 solute carrier family 19 (thiamine transporter), member 2 /FL=gb:AF153330.1 gb:AF135488.1 gb:AF160812.1	AF153330		NP_008927
209682_at	5.03E-04	gb:U26710.1 /DEF=Human cbl-b mRNA, complete cds. /FEA=mRNA /PROD=cbl-b /DB_XREF=gi:862406 /UG=Hs.3144 Cas-Br-M (murine) ectropic retroviral transforming sequence b /FL=gb:U26710.1	U26710		NP_733762

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209685_s_at	0.00228	gb:M13975.1 /DEF=Homo sapiens protein kinase C beta-II type (PRKCB1) mRNA, complete cds. /FEA=mRNA /GEN=PRKCB1 /PROD=protein kinase C beta-II type /DB_XREF=gi:189968 /UG=Hs.77202 protein kinase C, beta 1 /FL=gb:M13975.1	M13975		NP_002729
209688_s_at	0.00228	gb:BC005078.1 /DEF=Homo sapiens, clone MGC:13033, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:13033) /DB_XREF=gi:13477224 /UG=Hs.26118 Homo sapiens clone 24766 mRNA sequence /FL=gb:BC005078.1	BC005078		NP_113635
209689_at	0.011788	gb:BC005078.1 /DEF=Homo sapiens, clone MGC:13033, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:13033) /DB_XREF=gi:13477224 /UG=Hs.26118 Homo sapiens clone 24766 mRNA sequence /FL=gb:BC005078.1	BC005078		NP_113635
209694_at	0.045608	gb:M97655.1 /DEF=Human 6-pyruvoyltetrahydropterin synthase (PTS) mRNA, complete cds. /FEA=mRNA /GEN=PTS /PROD=6-pyruvoyltetrahydropterin synthase /DB_XREF=gi:306438 /UG=Hs.366 6-pyruvoyltetrahydropterin synthase /FL=gb:M97655.1 gb:NM_000317.1 gb:D17400.1	M97655		NP_000308
209707_at	1.26E-04	gb:AF022913.1 /DEF=Homo sapiens GPI transamidase mRNA, complete cds. /FEA=mRNA /PROD=GPI transamidase /DB_XREF=gi:2558890 /UG=Hs.62187 phosphatidylinositol glycan, class K /FL=gb:AF022913.1	AF022913		NP_005473
209711_at	0.003171	UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter	N80922	Hs.82635	NP_055954

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209732_at	0.02145	gb:BC005254.1 /DEF=Homo sapiens, Similar to C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced), clone MGC:12289, mRNA, complete cds. /FEA=mRNA /PROD=Similar to C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced) /DB_XREF=gi:13528920 /UG=Hs.85201 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced) /FL=gb:BC005254.1 gb:AB015628.1 gb:NM_005127.1	BC005254		NP_005118
209734_at	2.05E-04	gb:BC001604.1 /DEF=Homo sapiens, hematopoietic protein 1, clone MGC:2204, mRNA, complete cds. /FEA=mRNA /PROD=hematopoietic protein 1 /DB_XREF=gi:12804404 /UG=Hs.132834 hematopoietic protein 1 /FL=gb:BC001604.1 gb:M58285.1 gb:NM_005337.1	BC001604		NP_005328
209738_x_at	0.022758	gb:M31125.1 /DEF=Human pregnancy-specific-beta 1-glycoprotein (PS1) mRNA, complete cds, clone hPS12. /FEA=mRNA /DB_XREF=gi:190619 /UG=Hs.252097 pregnancy specific beta-1-glycoprotein 6 /FL=gb:M31125.1	M31125		NP_002773
209748_at	0.026891	gb:AB029006.1 /DEF=Homo sapiens mRNA for KIAA1083 protein, complete cds. /FEA=mRNA /GEN=KIAA1083 /PROD=KIAA1083 protein /DB_XREF=gi:5689502 /UG=Hs.26334 spastic paraplegia 4 (autosomal dominant; spastin) /FL=gb:AB029006.1	AB029006		NP_055761
209753_s_at	0.001116	thymopoietin	BG391171	Hs.11355	NP_003267

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209754_s_at	0.011626	gb:AF113682.1 /DEF=Homo sapiens clone FLB3436 PRO0868 mRNA, complete cds. /FEA=mRNA /PROD=PRO0868 /DB_XREF=gi:6855608 /UG=Hs.11355 thymopoietin /FL=gb:U09087.1 gb:AF113682.1	AF113682		NP_003267
209760_at	1.26E-04	gb:AL136932.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H1322 (from clone DKFZp586H1322); complete cds. /FEA=mRNA /GEN=DKFZp586H1322 /PROD=hypothetical protein /DB_XREF=gi:12053358 /UG=Hs.37892 KIAA0922 protein /FL=gb:AL136932.1	AL136932		NP_056011
209761_s_at	0.004341	Consensus includes gb:AA969194 /FEA=EST /DB_XREF=gi:3144374 /DB_XREF=est:op51c02.s1 /CLONE=IMAGE:1580354 /UG=Hs.38125 interferon-induced protein 75, 52kD /FL=gb:AF280094.1	AF280094		NP_536349
209771_x_at	1.26E-04	CD24 antigen (small cell lung carcinoma cluster 4 antigen)	AA761181	Hs.381003	NP_037362
209772_s_at	0.004659	gb:X69397.1 /DEF=H.sapiens CD24 gene, complete CDS. /FEA=mRNA /GEN=CD24 /PROD=cell surface antigen /DB_XREF=gi:396167 /UG=Hs.286124 CD24 antigen (small cell lung carcinoma cluster 4 antigen) /FL=gb:X69397.1	X69397		NP_037362
209773_s_at	1.58E-04	gb:BC001886.1 /DEF=Homo sapiens, ribonucleotide reductase M2 polypeptide, clone MGC:1551, mRNA, complete cds. /FEA=mRNA /PROD=ribonucleotide reductase M2 polypeptide /DB_XREF=gi:12804874 /UG=Hs.75319 ribonucleotide reductase M2 polypeptide /FL=gb:BC001886.1	BC001886		NP_001025

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209788_s_at	0.001618	gb:AF183569.1 /DEF=Homo sapiens aminopeptidase PILS (APPILS) mRNA, complete cds. /FEA=mRNA /GEN=APPILS /PROD=aminopeptidase PILS /DB_XREF=gi:6642986 /UG=Hs.247043 type 1 tumor necrosis factor receptor shedding aminopeptidase regulator /FL=gb:AF183569.1 gb:AF222340.1 gb:NM_016442.1	AF183569		NP_057526
209799_at	0.001109	gb:AF100763.1 /DEF=Homo sapiens AMP-activated kinase alpha 1 subunit mRNA, complete cds. /FEA=mRNA /PROD=AMP-activated kinase alpha 1 subunit /DB_XREF=gi:5410311 /UG=Hs.9247 protein kinase, AMP-activated, alpha 1 catalytic subunit /FL=gb:AB022017.1 gb:AF100763.1 gb:NM_006251.1	AF100763		NP_006242
209806_at	1.26E-04	gb:BC000893.1 /DEF=Homo sapiens, H2B histone family, member A, clone MGC:5132, mRNA, complete cds. /FEA=mRNA /PROD=H2B histone family, member A /DB_XREF=gi:12654150 /UG=Hs.247817 H2B histone family, member A /FL=gb:BC000893.1	BC000893		NP_542160
209819_at	0.041795	gb:AF241831.1 /DEF=Homo sapiens intracellular hyaluronan-binding protein mRNA, complete cds. /FEA=mRNA /PROD=intracellular hyaluronan-binding protein /DB_XREF=gi:7417371 /UG=Hs.301839 intracellular antigen detected by monoclonal antibody Ki-1; intracellular hyaluronan-binding protein /FL=gb:AF241831.1	AF241831		NP_055097

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209829_at	0.001116	gb:AB002384.1 /DEF=Human mRNA for KIAA0386 gene, complete cds. /FEA=mRNA /GEN=KIAA0386 /DB_XREF=gi:2224712 /UG=Hs.101359 chromosome 6 open reading frame 32 /FL=gb:AB002384.1	AB002384		NP_056948
209831_x_at	0.001198	gb:AB004574.1 /DEF=Homo sapiens mRNA for deoxyribonuclease II, complete cds. /FEA=mRNA /GEN=DNASE2 /PROD=deoxyribonuclease II /DB_XREF=gi:3184394 /UG=Hs.118243 deoxyribonuclease II, lysosomal /FL=gb:AF045937.1 gb:AF047016.1 gb:AB004574.1 gb:AF060222.1 gb:NM_001375.1	AB004574		NP_001366
209833_at	1.26E-04	gb:U79115.1 /DEF=Human death adaptor molecule RAIDD (RAIDD) mRNA, complete cds. /FEA=mRNA /GEN=RAIDD /PROD=death adaptor molecule RAIDD /DB_XREF=gi:1785556 /UG=Hs.155566 CASP2 and RIPK1 domain containing adaptor with death domain /FL=gb:U79115.1 gb:U84388.1 gb:NM_003805.1	U79115		NP_003796
209845_at	0.00161	gb:AF117233.1 /DEF=Homo sapiens znf-xp protein mRNA, complete cds. /FEA=mRNA /PROD=znf-xp protein /DB_XREF=gi:6563239 /UG=Hs.7838 makorin, ring finger protein, 1 /FL=gb:AF117233.1	AF117233		NP_038474

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209846_s_at	2.05E-04	gb:BC002832.1 /DEF=Homo sapiens, Similar to butyrophilin, subfamily 3, member A2, clone MGC:3790, mRNA, complete cds. /FEA=mRNA /PROD=Similar to butyrophilin, subfamily 3, member A2 /DB_XREF=gi:12803966 /UG=Hs.87497 butyrophilin, subfamily 3, member A2 /FL=gb:BC002832.1 gb:U90144.1 gb:U90546.1 gb:NM_007047.1	BC002832		NP_008978
209852_x_at	1.26E-04	gb:BC001423.1 /DEF=Homo sapiens, Similar to proteasome (prosome, macropain) 28 subunit, 3, clone MGC:1394, mRNA, complete cds. /FEA=mRNA /PROD=Similar to proteasome (prosome, macropain) 28subunit, 3 /DB_XREF=gi:12655138 /UG=Hs.152978 proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki) /FL=gb:BC001423.1 gb:BC002684.1	BC001423		NP_789839
209853_s_at	0.003741	gb:BC002684.1 /DEF=Homo sapiens, proteasome (prosome, macropain) 28 subunit, 3, clone MGC:3572, mRNA, complete cds. /FEA=mRNA /PROD=proteasome (prosome, macropain) 28 subunit, 3 /DB_XREF=gi:12803696 /UG=Hs.152978 proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki) /FL=gb:BC001423.1 gb:BC002684.1	BC002684		NP_789839

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209858_x_at	0.003171	gb:BC002877.1 /DEF=Homo sapiens, Similar to hypothetical protein FLJ11585, clone MGC:11258, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein FLJ11585 /DB_XREF=gi:12804048 /UG=Hs.315367 Homo sapiens, Similar to hypothetical protein FLJ11585, clone MGC:11258, mRNA, complete cds /FL=gb:BC002877.1	BC002877		NP_075563
209860_s_at	0.001116	gb:J04543.1 /DEF=Human synexin mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:338243 /UG=Hs.78637 annexin A7 /FL=gb:BC002632.1 gb:J04543.1 gb:NM_001156.2	J04543		NP_004025
209864_at	0.004341	gb:AB045118.1 /DEF=Homo sapiens FRAT2 mRNA, complete cds. /FEA=mRNA /GEN=FRAT2 /PROD=FRAT2 /DB_XREF=gi:13365650 /UG=Hs.140720 GSK-3 binding protein FRAT2 /FL=gb:AB045118.1	AB045118		NP_036215
209868_s_at	1.26E-04	gb:D28482.1 /DEF=Human scr2 mRNA for RNA binding protein SCR2, complete cds. /FEA=mRNA /GEN=scr2 /PROD=SCR2 /DB_XREF=gi:520588 /UG=Hs.241567 RNA binding motif, single stranded interacting protein 1 /FL=gb:D28482.1 gb:NM_016836.1 gb:NM_016838.1 gb:NM_002897.2	D28482		NP_058523

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209882_at	3.26E-04	gb:AF084462.1 /DEF=Homo sapiens GTP-binding protein ROC1 (ROC1) mRNA, complete cds. /FEA=mRNA /GEN=ROC1 /PROD=GTP-binding protein ROC1 /DB_XREF=gi:4234917 /UG=Hs.96038 Ric (Drosophila)-like, expressed in many tissues /FL=gb:U71203.1 gb:U78165.1 gb:AF084462.1 gb:NM_006912.1	AF084462		NP_008843
209892_at	1.72E-04	Consensus includes gb:AF305083.1 /DEF=Homo sapiens alpha(1,3)-fucosyltransferase IV (FUTIV) gene, 3 UTR. /FEA=mRNA /DB_XREF=gi:11096240 /UG=Hs.2173 fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific) /FL=gb:M58596.1 gb:M58597.1 gb:NM_002033.1	AF305083		NP_002024
209894_at	0.006698	gb:U50748.1 /DEF=Homo sapiens leptin receptor short form (db) mRNA, complete cds. /FEA=mRNA /GEN=db /PROD=leptin receptor /DB_XREF=gi:3236285 /UG=Hs.226627 leptin receptor /FL=gb:U50748.1	U50748		NP_002294
209896_s_at	0.02145	gb:AF119855.1 /DEF=Homo sapiens PRO1847 mRNA, complete cds. /FEA=mRNA /PROD=PRO1847 /DB_XREF=gi:7770146 /UG=Hs.285196 hypothetical protein PRO1847 /FL=gb:AF119855.1	AF119855		
209901_x_at	7.58E-04	gb:U19713.1 /DEF=Human allograft-inflammatory factor-1 mRNA, complete cds. /FEA=mRNA /PROD=allograft-inflammatory factor-1 /DB_XREF=gi:1122908 /UG=Hs.76364 allograft inflammatory factor 1 /FL=gb:U19713.1 gb:U49392.1 gb:D86438.1 gb:NM_001623.2	U19713		NP_116573

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209906_at	2.05E-04	gb:U62027.1 /DEF=Human anaphylatoxin C3a receptor (HNFAG09) mRNA, complete cds. /FEA=mRNA /GEN=HNFAG09 /PROD=anaphylatoxin C3a receptor /DB_XREF=gi:1511643 /UG=Hs.155935 complement component 3a receptor 1 /FL=gb:U28488.1 gb:U62027.1 gb:NM_004054.1	U62027		NP_004045
209907_s_at	0.00228	gb:AF182198.1 /DEF=Homo sapiens intersectin 2 long isoform (ITSN2) mRNA, complete cds. /FEA=mRNA /GEN=ITSN2 /PROD=intersectin 2 long isoform /DB_XREF=gi:7329075 /UG=Hs.166184 intersectin 2 /FL=gb:AF182198.1	AF182198		NP_671494
209911_x_at	0.001116	gb:BC002842.1 /DEF=Homo sapiens, H2B histone family, member B, clone MGC:3802, mRNA, complete cds. /FEA=mRNA /PROD=H2B histone family, member B /DB_XREF=gi:12803984 /UG=Hs.180779 H2B histone family, member B /FL=gb:NM_021063.1 gb:BC002842.1	BC002842		NP_619790
209916_at	3.56E-04	gb:BC002477.1 /DEF=Homo sapiens, clone MGC:3090, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3090) /DB_XREF=gi:12803318 /UG=Hs.271586 hypothetical protein DKFZp762M115 /FL=gb:BC002477.1	BC002477		NP_061176
209919_x_at	2.05E-04	gb:L20490.1 /DEF=Human gamma-glutamyl transpeptidase mRNA, complete cds. /FEA=mRNA /PROD=gamma-glutamyl transpeptidase /DB_XREF=gi:306748 /UG=Hs.284380 gamma-glutamyltransferase 1 /FL=gb:J05235.1 gb:L20490.1	L20490		NP_543029

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209930_s_at	0.026891	gb:L13974.1 /DEF=Human leucine zipper mRNA, complete cds. /FEA=mRNA /PROD=leucine zipper protein /DB_XREF=gi:506817 /UG=Hs.75643 nuclear factor (erythroid-derived 2), 45kD /FL=gb:BC005044.1 gb:L24122.1 gb:L13974.1	L13974		NP_006154
209943_at	0.00161	gb:AF176699.1 /DEF=Homo sapiens F-box protein FBL4 mRNA, complete cds. /FEA=mRNA /PROD=F-box protein FBL4 /DB_XREF=gi:6103636 /UG=Hs.49526 f-box and leucine-rich repeat protein 4 /FL=gb:AF176699.1 gb:AF199355.1 gb:NM_012160.1	AF176699		NP_036292
209944_at	3.26E-04	gb:BC000330.1 /DEF=Homo sapiens, Similar to clones 23667 and 23775 zinc finger protein, clone MGC:8520, mRNA, complete cds. /FEA=mRNA /PROD=Similar to clones 23667 and 23775 zinc fingerprotein /DB_XREF=gi:12653126 /UG=Hs.7137 clones 23667 and 23775 zinc finger protein /FL=gb:BC000330.1 gb:BC004357.1	BC000330		NP_067011
209960_at	0.001053	Consensus includes gb:X16323.1 /DEF=Human mRNA for hepatocyte growth factor (HGF). /FEA=mRNA /DB_XREF=gi:32081 /UG=Hs.809 hepatocyte growth factor (hepapoietin A; scatter factor) /FL=gb:M60718.1 gb:M29145.1 gb:M73239.1 gb:M73240.1	X16323		NP_000592

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
209964_s_at	0.007779	gb:AF032105.1 /DEF=Homo sapiens ataxin-7 (SCA7) mRNA, complete cds. /FEA=mRNA /GEN=SCA7 /PROD=ataxin-7 /DB_XREF=gi:3192953 /UG=Hs.108447 spinocerebellar ataxia 7 (olivopontocerebellar atrophy with retinal degeneration) /FL=gb:AF032105.1	AF032105		NP_000324
209984_at	0.003171	gb:AB037901.1 /DEF=Homo sapiens GASC-1 mRNA, complete cds. /FEA=mRNA /GEN=GASC-1 /DB_XREF=gi:10567163 /UG=Hs.149918 gene amplified in squamous cell carcinoma 1; KIAA0780 protein /FL=gb:AB037901.1	AB037901		NP_055876
209992_at	7.22E-04	gb:AB044805.1 /DEF=Homo sapiens mRNA for 6-phosphofructo-2-kinase heart isoform, complete cds. /FEA=mRNA /PROD=6-phosphofructo-2-kinase heart isoform /DB_XREF=gi:11933148 /UG=Hs.211585 6-phosphofructo-2-kinasefructose-2,6-biphosphatase 2 /FL=gb:AB044805.1	AB044805		NP_006203
209993_at	0.023827	gb:AF016535.1 /DEF=Homo sapiens P-glycoprotein (mdr1) mRNA, complete cds. /FEA=mRNA /GEN=mdr1 /PROD=P-glycoprotein /DB_XREF=gi:2353263 /UG=Hs.21330 ATP-binding cassette, sub-family B (MDRTAP), member 1 /FL=gb:M14758.1 gb:AF016535.1 gb:NM_000927.2	AF016535		NP_000918

G ne Identifier	p-value	Description	G ne Accession No.	Unigene Accession No.	Protein Accession No.
209997_x_at	0.009652	gb:BC000453.1 /DEF=Homo sapiens, Similar to pericentriolar material 1, clone MGC:8458, mRNA, complete cds. /FEA=mRNA /PROD=Similar to pericentriolar material 1 /DB_XREF=gi:12653366 /UG=Hs.315478 Homo sapiens, Similar to pericentriolar material 1, clone MGC:8458, mRNA, complete cds /FL=gb:BC000453.1	BC000453		NP_006188
210004_at	1.28E-04	gb:AF035776.1 /DEF=Homo sapiens oxidized low-density lipoprotein receptor mRNA, complete cds. /FEA=mRNA /PROD=oxidized low-density lipoprotein receptor /DB_XREF=gi:3941299 /UG=Hs.77729 oxidised low density lipoprotein (lectin-like) receptor 1 /FL=gb:AB010710.1 gb:AF035776.1 gb:NM_002543.1	AF035776		NP_002534
210009_s_at	0.034317	gb:AF229796.1 /DEF=Homo sapiens golgi SNARE isoform B (GOSR2) mRNA, complete cds, alternatively spliced. /FEA=mRNA /GEN=GOSR2 /PROD=golgi SNARE isoform B /DB_XREF=gi:12711466 /UG=Hs.100651 golgi SNAP receptor complex member 2 /FL=gb:AF229796.1 gb:AF007548.1 gb:NM_004287.1	AF229796		NP_473363
210014_x_at	0.033381	gb:AF023266.1 /DEF=Homo sapiens NAD ⁺ -specific isocitrate dehydrogenase beta subunit isoform B mRNA, nuclear gene encoding mitochondrial product, complete cds. /FEA=mRNA /PROD=NAD ⁺ -specific isocitrate dehydrogenase betasubunit isoform B /DB_XREF=gi:4103447 /UG=Hs.155410 isocitrate dehydrogenase 3 (NAD ⁺) beta /FL=gb:AF023266.1	AF023266		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210018_x_at	0.035611	gb:AB026118.1 /DEF=Homo sapiens mRNA for MALT1, complete cds. /FEA=mRNA /GEN=MALT1 /PROD=MALT1 /DB_XREF=gi:5706377 /UG=Hs.180566 mucosa associated lymphoid tissue lymphoma translocation gene 1 /FL=gb:AB026118.1	AB026118		NP_776216
210031_at	0.001054	gb:J04132.1 /DEF=Human T cell receptor zeta-chain mRNA, complete cds. /FEA=mRNA /PROD=T-cell receptor zeta chain /DB_XREF=gi:623041 /UG=Hs.97087 CD3Z antigen, zeta polypeptide (TiT3 complex) /FL=gb:NM_000734.1 gb:J04132.1	J04132		NP_000725
210044_s_at	0.004026	gb:BC002796.1 /DEF=Homo sapiens, lymphoblastic leukemia derived sequence 1, clone MGC:3943, mRNA, complete cds. /FEA=mRNA /PROD=lymphoblastic leukemia derived sequence 1 /DB_XREF=gi:12803898 /UG=Hs.46446 lymphoblastic leukemia derived sequence 1 /FL=gb:BC002796.1 gb:NM_005583.1	BC002796		NP_005574
210057_at	0.022708	gb:U32581.2 /DEF=Homo sapiens lambdaiota protein kinase C-interacting protein mRNA, complete cds. /FEA=mRNA /PROD=lambdaiota protein kinase C-interactingprotein /DB_XREF=gi:5542015 /UG=Hs.168052 KIAA0421 protein /FL=gb:U32581.2	U32581		NP_055907
210075_at	1.41E-04	gb:AF151074.1 /DEF=Homo sapiens HSPC240 mRNA, complete cds. /FEA=mRNA /PROD=HSPC240 /DB_XREF=gi:7106869 /UG=Hs.132744 hypothetical protein /FL=gb:AF151074.1 gb:NM_016496.1	AF151074		NP_057580

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210076_x_at	0.02145	gb:AF151813.1 /DEF=Homo sapiens CGI-55 protein mRNA, complete cds. /FEA=mRNA /PROD=CGI-55 protein /DB_XREF=gi:4929578 /UG=Hs.165998 PAI-1 mRNA-binding protein /FL=gb:AF151813.1	AF151813		NP_056455
210077_s_at	0.010061	gb:U30884.1 /DEF=Human splicing factor SRp40-2 (SRp40) mRNA, complete cds. /FEA=mRNA /GEN=SRp40 /PROD=SRp40-2 /DB_XREF=gi:1049089 /UG=Hs.166975 splicing factor, arginineserine-rich 5 /FL=gb:U30884.1	U30884		NP_008856
210088_x_at	2.17E-04	gb:M36172.1 /DEF=Human embryonic myosin alkali light chain (MLC1) mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:188580 /UG=Hs.298161 myosin, light polypeptide 4, alkali; atrial, embryonic /FL=gb:M36172.1 gb:M24121.1 gb:NM_002476.1	M36172		NP_002467
210092_at	0.02145	gb:AF067173.1 /DEF=Homo sapiens Mago homolog mRNA, complete cds. /FEA=mRNA /PROD=Mago homolog /DB_XREF=gi:4894379 /UG=Hs.57904 mago-nashi (Drosophila) homolog, proliferation associated /FL=gb:AF035940.1 gb:AF067173.1 gb:NM_002370.2	AF067173		NP_002361
210097_s_at	0.033381	Consensus includes gb:AF130102.1 /DEF=Homo sapiens clone FLB2205 PRO0522 mRNA, complete cds. /FEA=mRNA /PROD=PRO0522 /DB_XREF=gi:11493507 /UG=Hs.106346 retinoic acid repressible protein /FL=gb:AF130102.1	AF130102		NP_057251

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210101_x_at	1.26E-04	gb:AF257318.1 /DEF=Homo sapiens SH3-containing protein SH3GLB1 mRNA, complete cds. /FEA=mRNA /PROD=SH3-containing protein SH3GLB1 /DB_XREF=gi:8896091 /UG=Hs.136309 SH3-containing protein SH3GLB1 /FL=gb:AF350371.1 gb:AF151819.1 gb:NM_016009.1 gb:AF257318.1	AF257318		NP_057093
210102_at	0.021045	gb:BC001234.1 /DEF=Homo sapiens, Similar to loss of heterozygosity, 11, chromosomal region 2, gene A, clone MGC:4904, mRNA, complete cds. /FEA=mRNA /PROD=Similar to loss of heterozygosity, 11,chromosomal region 2, gene A /DB_XREF=gi:12654782 /UG=Hs.152944 loss of heterozygosity, 11, chromosomal region 2, gene A /FL=gb:BC001234.1	BC001234		NP_055437
210113_s_at	0.003539	gb:AF310105.1 /DEF=Homo sapiens NALP1 mRNA, complete cds. /FEA=mRNA /PROD=NALP1 /DB_XREF=gi:11096302 /UG=Hs.104305 death effector filament-forming Ced-4-like apoptosis protein /FL=gb:AF310105.1 gb:AF229059.1 gb:AB023143.1 gb:NM_014922.1	AF310105		NP_127500
210117_at	0.00161	gb:AF311312.1 /DEF=Homo sapiens infertility-related sperm protein mRNA, complete cds. /FEA=mRNA /PROD=infertility-related sperm protein /DB_XREF=gi:10863767 /UG=Hs.153057 sperm associated antigen 1 /FL=gb:AF311312.1 gb:NM_003114.1	AF311312		NP_757367

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210119_at	3.26E-04	gb:U73191.1 /DEF=Human inward rectifier potassium channel (Kir1.3), complete cds. /FEA=mRNA /GEN=Kir1.3 /PROD=inward rectifier potassium channel /DB_XREF=gi:1765984 /UG=Hs.17287 potassium inwardly-rectifying channel, subfamily J, member 15 /FL=gb:U73191.1 gb:NM_002243.1	U73191		NP_733933
210128_s_at	0.003862	gb:U41070.1 /DEF=Human P2 purinergic receptor mRNA, complete cds. /FEA=mRNA /PROD=P2 purinergic receptor /DB_XREF=gi:1469913 /UG=Hs.28408 leukotriene b4 receptor (chemokine receptor-like 1) /FL=gb:U41070.1 gb:D89079.1	U41070		NP_000743
210131_x_at	3.26E-04	gb:D49737.1 /DEF=Homo sapiens mRNA for cytochrome b large subunit of complex II, complete cds. /FEA=mRNA /PROD=cytochrome b large subunit of complex II /DB_XREF=gi:2588778 /UG=Hs.3577 succinate dehydrogenase complex, subunit C, integral membrane protein, 15kD /FL=gb:D49737.1	D49737		NP_002992
210136_at	5.03E-04	ESTs, Moderately similar to MBP_HUMAN Myelin basic protein (MBP) (Myelin A1 protein) (Myelin membrane encephalitogenic protein) [H.sapiens]	AW070431		NP_002376
210140_at	0.00161	gb:AF031824.1 /DEF=Homo sapiens leukocystatin mRNA, complete cds. /FEA=mRNA /PROD=leukocystatin /DB_XREF=gi:3252857 /UG=Hs.143212 cystatin F (leukocystatin) /FL=gb:AF031824.1 gb:AF036342.1 gb:AB015225.1 gb:NM_003650.1	AF031824	Hs.409352	NP_003641

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210142_x_at	2.05E-04	gb:AF117234.1 /DEF=Homo sapiens flotillin mRNA, complete cds. /FEA=mRNA /PROD=flotillin /DB_XREF=gi:6563241 /UG=Hs.179986 flotillin 1 /FL=gb:AF117234.1	AF117234		NP_005794
210144_at	0.021165	Consensus includes gb:AK000851.1 /DEF=Homo sapiens cDNA FLJ20844 fis, clone ADKA01904. /FEA=mRNA /DB_XREF=gi:7021173 /UG=Hs.20017 chromosome 22 open reading frame 4 /FL=gb:BC002743.1	AK000851		NP_055161
210145_at	0.00161	gb:M68874.1 /DEF=Homo sapiens phosphatidylcholine 2-acylhydrolase (cPLA2) mRNA, complete cds. /FEA=mRNA /GEN=cPLA2 /PROD=phosphatidylcholine 2-acylhydrolase /DB_XREF=gi:190003 /UG=Hs.211587 phospholipase A2, group IVA (cytosolic, calcium-dependent) /FL=gb:M68874.1 gb:M72393.1	M68874		NP_077734
210146_x_at	0.016934	gb:AF004231.1 /DEF=Homo sapiens monocytemacrophage Ig-related receptor MIR-10 (MIR cl-10) mRNA, complete cds. /FEA=mRNA /GEN=MIR cl-10 /PROD=MIR-10 /DB_XREF=gi:2343110 /UG=Hs.22405 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2 /FL=gb:AF000574.1 gb:AF004231.1 gb:AF011565.1	AF004231		NP_005865
210149_s_at	1.26E-04	gb:AF061735.1 /DEF=Homo sapiens clone 023e08 My032 protein mRNA, complete cds. /FEA=mRNA /PROD=My032 protein /DB_XREF=gi:12002005 /UG=Hs.307099 Homo sapiens clone 023e08 My032 protein mRNA, complete cds /FL=gb:AF061735.1	AF061735		NP_006347

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210153_s_at	2.05E-04	gb:M55905.1 /DEF=Human mitochondrial NAD(P)+ dependent malic enzyme mRNA, complete cds. /FEA=mRNA /GEN=NAD(P)+ -dependent malic enzyme /PROD=mitochondrial NAD(P)+ -dependent malic enzyme /DB_XREF=gi:187299 /UG=Hs.75342 malic enzyme 2, NAD(+)-dependent, mitochondrial /FL=gb:NM_002396.2 gb:M55905.1	M55905		NP_002387
210154_at	5.64E-04	gb:M55905.1 /DEF=Human mitochondrial NAD(P)+ dependent malic enzyme mRNA, complete cds. /FEA=mRNA /GEN=NAD(P)+ -dependent malic enzyme /PROD=mitochondrial NAD(P)+ -dependent malic enzyme /DB_XREF=gi:187299 /UG=Hs.75342 malic enzyme 2, NAD(+)-dependent, mitochondrial /FL=gb:NM_002396.2 gb:M55905.1	M55905		NP_002387
210156_s_at	1.26E-04	gb:D25547.1 /DEF=Homo sapiens mRNA for PIMT isozyme I, complete cds. /FEA=mRNA /PROD=PIMT isozyme I /DB_XREF=gi:1332402 /UG=Hs.79137 protein-L-isoaspartate (D-aspartate) O-methyltransferase /FL=gb:D25547.1	D25547		NP_005380
210162_s_at	0.041055	gb:U08015.1 /DEF=Human NF-ATc mRNA, complete cds. /FEA=mRNA /PROD=NF-ATc /DB_XREF=gi:500631 /UG=Hs.96149 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1 /FL=gb:U08015.1	U08015		NP_765978

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210176_at	0.003171	gb:AL050262.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564I0682 (from clone DKFZp564I0682); complete cds. /FEA=mRNA /GEN=DKFZp564I0682 /PROD=hypothetical protein /DB_XREF=gi:4886482 /UG=Hs.2474 toll-like receptor 1 /FL=gb:U88540.1 gb:D13637.1 gb:NM_003263.1 gb:AL050262.1	AL050262		NP_003254
210187_at	0.02145	gb:BC005147.1 /DEF=Homo sapiens, FK506-binding protein 1A (12kD), clone MGC:2167, mRNA, complete cds. /FEA=mRNA /PROD=FK506-binding protein 1A (12kD) /DB_XREF=gi:13477342 /UG=Hs.752 FK506-binding protein 1A (12kD) /FL=gb:BC005147.1	BC005147		NP_463460
210191_s_at	1.26E-04	gb:BC002447.1 /DEF=Homo sapiens, Similar to putative homeodomain transcription factor, clone MGC:1307, mRNA, complete cds. /FEA=mRNA /PROD=Similar to putative homeodomain transcriptionfactor /DB_XREF=gi:12803264 /UG=Hs.123637 putative homeodomain transcription factor /FL=gb:BC002447.1	BC002447		NP_006599
210200_at	7.58E-04	gb:BC000108.1 /DEF=Homo sapiens, Similar to Nedd-4-like ubiquitin-protein ligase, clone MGC:2079, mRNA, complete cds. /FEA=mRNA /PROD=Similar to Nedd-4-like ubiquitin-protein ligase /DB_XREF=gi:12652716 /UG=Hs.19301 Homo sapiens, Similar to Nedd-4-like ubiquitin-protein ligase, clone MGC:2079, mRNA, complete cds /FL=gb:BC000108.1	BC000108		NP_008945

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210205_at	0.010205	gb:AB026730.1 /DEF=Homo sapiens B3GALT4 mRNA for beta-1,3-galactosyltransferase-4, complete cds. /FEA=mRNA /GEN=B3GALT4 /PROD=beta-1,3-galactosyltransferase-4 /DB_XREF=gi:6683013 /UG=Hs.21495 UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 4 /FL=gb:AF063595.1 gb:NM_003782.1 gb:AB026730.1	AB026730		NP_003773
210213_s_at	5.03E-04	gb:AF022229.1 /DEF=Homo sapiens translation initiation factor 6 (eIF6) mRNA, complete cds. /FEA=mRNA /GEN=eIF6 /PROD=translation initiation factor 6 /DB_XREF=gi:2809382 /UG=Hs.5215 integrin beta 4 binding protein /FL=gb:AF022229.1	AF022229		NP_002203
210218_s_at	0.012818	gb:U36501.1 /DEF=Human SP100 B (SP100-B) mRNA, complete cds. /FEA=mRNA /GEN=SP100-B /PROD=SP100-B /DB_XREF=gi:1173655 /UG=Hs.77617 nuclear antigen Sp100 /FL=gb:U36501.1	U36501		NP_003104
210225_x_at	0.013223	gb:AF009635.1 /DEF=Homo sapiens clone 17.7 immunoglobulin-like transcript 5 protein mRNA, complete cds. /FEA=mRNA /PROD=immunoglobulin-like transcript 5 protein /DB_XREF=gi:2662429 /UG=Hs.105928 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3 /FL=gb:AF000575.1 gb:U91928.1 gb:AF025533.1 gb:AF009632.1 gb:AF009633.1 gb:AF009635.1 gb:AF009638.1 gb:AF009639.1 gb:AF009641.1 gb:AF009642.1 gb:AF031553.1 gb:AF031554.1 gb:AF031555.1 gb:AF031556.1 gb:NM_006864.1	AF009635		NP_006855

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210231_x_at	1.26E-04	gb:D45198.1 /DEF=Human mRNA for template acyivating factor-I alpha, complete cds. /FEA=mRNA /GEN=set /PROD=template acyivating factor-I alpha /DB_XREF=gi:971271 /UG=Hs.145279 SET translocation (myeloid leukemia-associated) /FL=gb:D45198.1	D45198		NP_003002
210240_s_at	0.003529	gb:U20498.1 /DEF=Human p19 protein mRNA, complete cds. /FEA=mRNA /PROD=p19 protein /DB_XREF=gi:1161921 /UG=Hs.29656 cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4) /FL=gb:U40343.1 gb:U20498.1 gb:BC001822.1 gb:U49399.1 gb:NM_001800.1	U20498		NP_524145
210241_s_at	0.027239	gb:AB007458.1 /DEF=Homo sapiens mRNA for P53TG1-D, complete cds. /FEA=mRNA /GEN=P53TG1 /PROD=P53TG1-D /DB_XREF=gi:5006272 /UG=Hs.306301 Homo sapiens mRNA for P53TG1-D, complete cds /FL=gb:AB007458.1	AB007458		NP_009164
210244_at	1.41E-04	gb:U19970.1 /DEF=Human antimicrobial LPS-binding protein CAP18 precursor mRNA, complete cds. /FEA=mRNA /PROD=CAP18 precursor /DB_XREF=gi:643476 /UG=Hs.51120 cathelicidin antimicrobial peptide /FL=gb:NM_004345.1 gb:U19970.1	U19970		NP_004336
210249_s_at	0.00228	gb:U59302.1 /DEF=Human steroid receptor coactivator-1 F-SRC-1 mRNA, complete cds. /FEA=mRNA /PROD=steroid receptor coactivator-1 F-SRC-1 /DB_XREF=gi:1480645 /UG=Hs.74002 nuclear receptor coactivator 1 /FL=gb:U40396.1 gb:U59302.1 gb:U90661.1 gb:NM_003743.1	U59302		NP_671766

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210254_at	1.41E-04	gb:L35848.1 /DEF=Homo sapiens IgE receptor beta chain (HTm4) mRNA, complete cds. /FEA=mRNA /GEN=HTm4 /PROD=IgE receptor beta subunit /DB_XREF=gi:561638 /UG=Hs.99960 membrane-spanning 4-domains, subfamily A, member 3 (hematopoietic cell-specific) /FL=gb:NM_006138.1 gb:L35848.1	L35848		NP_006129
210255_at	0.019999	gb:U84138.1 /DEF=Homo sapiens DNA repair protein RAD51B mRNA, complete cds. /FEA=mRNA /PROD=DNA repair protein RAD51B /DB_XREF=gi:2801404 /UG=Hs.100669 RAD51 (S. cerevisiae)-like 1 /FL=gb:NM_002877.1 gb:U92074.1 gb:U84138.1	U84138		NP_598194
210260_s_at	0.02145	gb:BC005352.1 /DEF=Homo sapiens, TNF-induced protein, clone MGC:12451, mRNA, complete cds. /FEA=mRNA /PROD=TNF-induced protein /DB_XREF=gi:13529163 /UG=Hs.17839 TNF-induced protein /FL=gb:BC005352.1 gb:AF099935.1	BC005352		NP_055165
210275_s_at	2.05E-04	gb:AF062347.1 /DEF=Homo sapiens zinc finger protein 216 splice variant 2 (ZNF216) mRNA, complete cds. /FEA=mRNA /GEN=ZNF216 /PROD=zinc finger protein 216 splice variant 2 /DB_XREF=gi:3643810 /UG=Hs.3776 zinc finger protein 216 /FL=gb:AF062347.1	AF062347		NP_005998
210279_at	0.033381	gb:AF261135.1 /DEF=Homo sapiens GPR18-iso mRNA, complete cds. /FEA=mRNA /PROD=GPR18-iso /DB_XREF=gi:12005919 /UG=Hs.88269 Homo sapiens clone IMAGE:1837189, mRNA sequence /FL=gb:AF261135.1	AF261135		NP_005283

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210283_x_at	7.58E-04	gb:BC005295.1 /DEF=Homo sapiens, Similar to polyadenylate binding protein-interacting protein 1, clone MGC:12360, mRNA, complete cds. /FEA=mRNA /PROD=Similar to polyadenylate bindingprotein-interacting protein 1 /DB_XREF=gi:13529010 /UG=Hs.109643 polyadenylate binding protein-interacting protein 1 /FL=gb:BC005295.1	BC005295		NP_006442
210284_s_at	0.041055	gb:AF241230.1 /DEF=Homo sapiens TAK1-binding protein 2 mRNA, complete cds. /FEA=mRNA /PROD=TAK1-binding protein 2 /DB_XREF=gi:7677465 /UG=Hs.109727 TAK1-binding protein 2; KIAA0733 protein /FL=gb:AF241230.1	AF241230		NP_663317
210285_x_at	0.013223	gb:BC000383.1 /DEF=Homo sapiens, Wilms tumour 1-associating protein, clone MGC:8419, mRNA, complete cds. /FEA=mRNA /PROD=Wilms tumour 1-associating protein /DB_XREF=gi:12653228 /UG=Hs.119 Wilms tumour 1-associating protein /FL=gb:BC000383.1 gb:BC004432.1	BC000383		NP_690597
210288_at	0.00161	gb:AF081675.1 /DEF=Homo sapiens ITIM-containing receptor MAFA-L mRNA, complete cds. /FEA=mRNA /PROD=ITIM-containing receptor MAFA-L /DB_XREF=gi:3421400 /UG=Hs.146322 killer cell lectin-like receptor subfamily G, member 1 /FL=gb:AF034952.1 gb:AF081675.1 gb:AF097358.1 gb:NM_005810.1	AF081675		NP_005801

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210293_s_at	1.26E-04	gb:BC005032.1 /DEF=Homo sapiens, Sec23 (S. cerevisiae) homolog B, clone MGC:12666, mRNA, complete cds. /FEA=mRNA /PROD=Sec23 (S. cerevisiae) homolog B /DB_XREF=gi:13477148 /UG=Hs.173497 Sec23 (S. cerevisiae) homolog B /FL=gb:BC005032.1	BC005032		NP_116781
210296_s_at	5.03E-04	gb:BC005375.1 /DEF=Homo sapiens, peroxisomal membrane protein 3 (35kD, Zellweger syndrome), clone MGC:12491, mRNA, complete cds. /FEA=mRNA /PROD=peroxisomal membrane protein 3 (35kD, Zellwegersyndrome) /DB_XREF=gi:13529226 /UG=Hs.180612 peroxisomal membrane protein 3 (35kD, Zellweger syndrome) /FL=gb:BC005375.1	BC005375		NP_000309
210299_s_at	7.57E-04	gb:AF063002.1 /DEF=Homo sapiens LIM protein SLIMMER mRNA, complete cds. /FEA=mRNA /PROD=LIM protein SLIMMER /DB_XREF=gi:3859848 /UG=Hs.239069 four and a half LIM domains 1 /FL=gb:AF098518.1 gb:AF063002.1	AF063002		NP_001440
210312_s_at	1.26E-04	gb:BC002640.1 /DEF=Homo sapiens, Similar to uterine protein, clone MGC:4279, mRNA, complete cds. /FEA=mRNA /PROD=Similar to uterine protein /DB_XREF=gi:12803610 /UG=Hs.4187 hypothetical protein 24636 /FL=gb:BC002640.1	BC002640		NP_777547

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210314_x_at	2.05E-04	gb:AF114013.1 /DEF=Homo sapiens tumor necrosis factor-related death ligand-1gamma mRNA, complete cds. /FEA=mRNA /PROD=tumor necrosis factor-related deathligand 1gamma /DB_XREF=gi:7328557 /UG=Hs.54673 tumor necrosis factor (ligand) superfamily, member 13 /FL=gb:AF136294.1 gb:AF114013.1	AF114013		NP_742086
210340_s_at	0.001116	gb:BC002635.1 /DEF=Homo sapiens, colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage), clone MGC:3848, mRNA, complete cds. /FEA=mRNA /PROD=colony stimulating factor 2 receptor, alpha,low-affinity (granulocyte-macrophage) /DB_XREF=gi:12803600 /UG=Hs.182378 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage) /FL=gb:BC002635.1	BC002635		NP_758452
210356_x_at	0.003539	gb:BC002807.1 /DEF=Homo sapiens, membrane-spanning 4-domains, subfamily A, member 2, clone MGC:3969, mRNA, complete cds. /FEA=mRNA /PROD=membrane-spanning 4-domains, subfamily A, member2 /DB_XREF=gi:12803920 /UG=Hs.89751 membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE, high affinity I, receptor for; beta polypeptide) /FL=gb:NM_021950.1 gb:BC002807.1	BC002807		NP_690606

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210360_s_at	0.037567	gb:AF116674.1 /DEF=Homo sapiens PRO1941 mRNA, complete cds. /FEA=mRNA /PROD=PRO1941 /DB_XREF=gi:7959846 /UG=Hs.77694 KIAA0429 gene product /FL=gb:AF116674.1	AF116674		NP_055566
210367_s_at	0.02145	gb:AF010316.1 /DEF=Homo sapiens Pig12 (PIG12) mRNA, complete cds. /FEA=mRNA /GEN=PIG12 /PROD=Pig12 /DB_XREF=gi:2415307 /UG=Hs.146688 prostaglandin E synthase /FL=gb:AF010316.1	AF010316		NP_004869
210386_s_at	7.58E-04	gb:BC001906.1 /DEF=Homo sapiens, Similar to metaxin 1, clone MGC:2518, mRNA, complete cds. /FEA=mRNA /PROD=Similar to metaxin 1 /DB_XREF=gi:12804906 /UG=Hs.247551 metaxin 1 /FL=gb:BC001906.1	BC001906		NP_002446
210395_x_at	3.26E-04	gb:AF116676.1 /DEF=Homo sapiens PRO1957 mRNA, complete cds. /FEA=mRNA /PROD=PRO1957 /DB_XREF=gi:7959850 /UG=Hs.298161 myosin, light polypeptide 4, alkali; atrial, embryonic /FL=gb:AF116676.1	AF116676		NP_002467
210401_at	0.005852	gb:U45448.1 /DEF=Human P2x1 receptor mRNA, complete cds. /FEA=mRNA /PROD=P2X1 receptor /DB_XREF=gi:1314356 /UG=Hs.41735 purinergic receptor P2X, ligand-gated ion channel, 1 /FL=gb:U45448.1 gb:AF020498.1 gb:NM_002558.1	U45448		NP_002549
210406_s_at	1.26E-04	gb:AL136727.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566K144 (from clone DKFZp566K144); complete cds. /FEA=mRNA /GEN=DKFZp566K144 /PROD=hypothetical protein /DB_XREF=gi:12052972 /UG=Hs.5636 RAB6A, member RAS oncogene family /FL=gb:AL136727.1	AL136727		NP_115520

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210407_at	0.013088	gb:AF070670.1 /DEF=Homo sapiens protein phosphatase 2C alpha 2 mRNA, complete cds. /FEA=mRNA /PROD=protein phosphatase 2C alpha 2 /DB_XREF=gi:3372871 /UG=Hs.57764 protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform /FL=gb:AF070670.1	AF070670		NP_808821
210422_x_at	0.02145	gb:D50402.1 /DEF=Human mRNA for NRAMP1, complete cds. /FEA=mRNA /PROD=Nramp /DB_XREF=gi:1000996 /UG=Hs.182611 solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1 /FL=gb:D50402.1 gb:D50403.1 gb:NM_000578.1 gb:L32185.1	D50402		NP_000569
210434_x_at	0.010205	gb:AF151056.1 /DEF=Homo sapiens HSPC222 mRNA, complete cds. /FEA=mRNA /PROD=HSPC222 /DB_XREF=gi:7106833 /UG=Hs.323093 Homo sapiens, jumping translocation breakpoint, clone MGC:10274, mRNA, complete cds /FL=gb:AF151056.1	AF151056		NP_006685
210438_x_at	0.016934	gb:M25077.1 /DEF=Human SS-Aro ribonucleoprotein autoantigen 60 kd subunit mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:387656 /UG=Hs.554 Sjogren syndrome antigen A2 (60kD, ribonucleoprotein autoantigen SS-Aro) /FL=gb:M25077.1	M25077		NP_004591

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
210453_x_at	5.03E-04	gb:AL050277.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566G013 (from clone DKFZp566G013); complete cds. /FEA=mRNA /GEN=DKFZp566G013 /PROD=hypothetical protein /DB_XREF=gi:4886454 /UG=Hs.107476 ATP synthase, H ⁺ transporting, mitochondrial F1F0, subunit g /FL=gb:AL050277.1	AL050277		NP_006467
210460_s_at	1.26E-04	gb:AB033605.1 /DEF=Homo sapiens mRNA for pUb-R5, complete cds. /FEA=mRNA /GEN=hpUb-R5 /PROD=pUb-R5 /DB_XREF=gi:8918352 /UG=Hs.148495 proteasome (prosome, macropain) 26S subunit, non-ATPase, 4 /FL=gb:AB033605.1	AB033605		NP_722544
210479_s_at	0.003539	gb:L14611.1 /DEF=Human transcription factor RZR-alpha mRNA, complete cds. /FEA=mRNA /PROD=transcription factor /DB_XREF=gi:348240 /UG=Hs.2156 RAR-related orphan receptor A /FL=gb:L14611.1 gb:NM_002943.1 gb:U04899.1	L14611		NP_599024

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210483_at	0.04734	gb:BC005043.1 /DEF=Homo sapiens, Similar to tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain, clone MGC:12787, mRNA, complete cds. /FEA=mRNA /PROD=Similar to tumor necrosis factor receptorsuperfamily, member 10c, decoy without an intracellulardomain /DB_XREF=gi:13477162 /UG=Hs.253829 Homo sapiens, Similar to tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain, clone MGC:12787, mRNA, complete cds /FL=gb:BC005043.1	BC005043		
210499_s_at	0.026891	gb:AB041834.1 /DEF=Homo sapiens mRNA for PQBP-1bc, complete cds. /FEA=mRNA /PROD=PQBP-1bc /DB_XREF=gi:10801589 /UG=Hs.30570 polyglutamine binding protein 1 /FL=gb:AB041834.1	AB041834		NP_005701
210504_at	0.001477	gb:U65404.1 /DEF=Human erythroid-specific transcription factor EKLF mRNA, complete cds. /FEA=mRNA /PROD=erythroid-specific transcription factor EKLF /DB_XREF=gi:1552995 /UG=Hs.37860 Kruppel-like factor 1 (erythroid) /FL=gb:NM_006563.1 gb:U65404.1	U65404		NP_006554
210506_at	0.003741	gb:U11282.1 /DEF=Human alpha(1,3)fucosyltransferase mRNA, complete cds. /FEA=mRNA /PROD=alpha(1,3)fucosyltransferase /DB_XREF=gi:520524 /UG=Hs.457 fucosyltransferase 7 (alpha (1,3) fucosyltransferase) /FL=gb:NM_004479.1 gb:U08112.1 gb:U11282.1	U11282		NP_004470

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210512_s_at	0.003862	gb:AF022375.1 /DEF=Homo sapiens vascular endothelial growth factor mRNA, complete cds. /FEA=mRNA /PROD=vascular endothelial growth factor /DB_XREF=gi:3719220 /UG=Hs.73793 vascular endothelial growth factor /FL=gb:M32977.1 gb:AF022375.1 gb:NM_003376.1 gb:AB021221.1 gb:AF091352.1	AF022375		NP_003367
210532_s_at	1.26E-04	gb:AF116639.1 /DEF=Homo sapiens PRO1574 mRNA, complete cds. /FEA=mRNA /PROD=PRO1574 /DB_XREF=gi:7959779 /UG=Hs.109052 chromosome 14 open reading frame 2 /FL=gb:AF116639.1	AF116639		
210542_s_at	0.004341	gb:BC000585.1 /DEF=Homo sapiens, Similar to solute carrier family 21 (organic anion transporter), member 11, clone MGC:659, mRNA, complete cds. /FEA=mRNA /PROD=Similar to solute carrier family 21 (organicanion transporter), member 11 /DB_XREF=gi:12653614 /UG=Hs.14805 solute carrier family 21 (organic anion transporter), member 11 /FL=gb:BC000585.1	BC000585		NP_037404
210556_at	0.0174	gb:U85430.1 /DEF=Human transcription factor NFATx4 mRNA, complete cds. /FEA=mRNA /PROD=transcription factor NFATx4 /DB_XREF=gi:1835590 /UG=Hs.172674 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3 /FL=gb:U85430.1	U85430		NP_775188

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210573_s_at	0.042536	gb:BC004424.1 /DEF=Homo sapiens, clone MGC:3538, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3538) /DB_XREF=gi:13325209 /UG=Hs.250745 polymerase (RNA) III (DNA directed) (62kD) /FL=gb:BC004424.1	BC004424		NP_006459
210582_s_at	0.016934	gb:AL117466.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586K0922 (from clone DKFZp586K0922); complete cds. /FEA=mRNA /GEN=DKFZp586K0922 /PROD=hypothetical protein /DB_XREF=gi:5911930 /UG=Hs.278027 LIM domain kinase 2 /FL=gb:AL117466.1 gb:NM_016733.1	AL117466		NP_057952
210592_s_at	0.033381	gb:M55580.1 /DEF=Human spermidinespermine N1-acetyltransferase mRNA, complete cds. /FEA=mRNA /GEN=spermidinespermine N1-acetyltransferase /PROD=spermidinespermine N1-acetyltransferase /DB_XREF=gi:338335 /UG=Hs.28491 spermidinespermine N1-acetyltransferase /FL=gb:M55580.1	M55580		
210598_at	0.004341	gb:AF130051.1 /DEF=Homo sapiens clone FLB3535 PRO0898 mRNA, complete cds. /FEA=mRNA /PROD=PRO0898 /DB_XREF=gi:11493408 /UG=Hs.306960 Homo sapiens clone FLB3535 PRO0898 mRNA, complete cds /FL=gb:AF130051.1	AF130051		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210608_s_at	0.017832	gb:BC001899.1 /DEF=Homo sapiens, Similar to fucosyltransferase 2 (secretor status included), clone MGC:2062, mRNA, complete cds. /FEA=mRNA /PROD=Similar to fucosyltransferase 2 (secretor statusincluded) /DB_XREF=gi:12804894 /UG=Hs.46328 fucosyltransferase 2 (secretor status included) /FL=gb:BC001899.1	BC001899		NP_000502
210621_s_at	0.003171	gb:M23612.1 /DEF=Human GTPase-activating protein (GAP) mRNA, complete cds. /FEA=mRNA /GEN=GAP /PROD=GTPase-activating protein /DB_XREF=gi:182973 /UG=Hs.758 RAS p21 protein activator (GTPase activating protein) 1 /FL=gb:NM_022650.1 gb:M23612.1	M23612		NP_072179
210624_s_at	0.010205	gb:BC000109.1 /DEF=Homo sapiens, Similar to ilvB (bacterial acetolactate synthase)-like, clone MGC:1269, mRNA, complete cds. /FEA=mRNA /PROD=Similar to ilvB (bacterial acetolactatesynthase)-like /DB_XREF=gi:12652718 /UG=Hs.78880 ilvB (bacterial acetolactate synthase)-like /FL=gb:BC000109.1	BC000109		NP_789796
210629_x_at	0.004341	gb:AF000425.1 /DEF=Homo sapiens LST1 mRNA, cLST1A splice variant, complete cds. /FEA=mRNA /GEN=LST1 /DB_XREF=gi:2145065 /UG=Hs.88411 lymphocyte antigen 117 /FL=gb:AF000425.1	AF000425		NP_009092
210638_s_at	1.26E-04	gb:AF176704.1 /DEF=Homo sapiens F-box protein FBX9 mRNA, complete cds. /FEA=mRNA /PROD=F-box protein FBX9 /DB_XREF=gi:6103646 /UG=Hs.11050 F-box only protein 9 /FL=gb:AF176704.1	AF176704		NP_258442

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210639_s_at	2.05E-04	gb:AF293841.1 /DEF=Homo sapiens apoptosis-related protein (APG5L) mRNA, complete cds, alternatively spliced. /FEA=mRNA /GEN=APG5L /PROD=apoptosis-related protein /DB_XREF=gi:12006863 /UG=Hs.11171 APG5 (autophagy 5, <i>S. cerevisiae</i>)-like /FL=gb:AF293841.1	AF293841		NP_004840
210644_s_at	7.58E-04	gb:AF109683.1 /DEF=Homo sapiens leukocyte-associated Ig-like receptor 1b mRNA, complete cds. /FEA=mRNA /PROD=leukocyte-associated Ig-like receptor 1b /DB_XREF=gi:6563041 /UG=Hs.115808 leukocyte-associated Ig-like receptor 1 /FL=gb:AF251509.2 gb:NM_021706.1 gb:AF109683.1	AF109683		NP_068354
210646_x_at	0.007779	gb:BC001675.1 /DEF=Homo sapiens, ribosomal protein L13a, clone MGC:2546, mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein L13a /DB_XREF=gi:12804528 /UG=Hs.119122 ribosomal protein L13a /FL=gb:BC001675.1 gb:BC001836.1	BC001675		NP_036555
210647_x_at	0.004341	gb:AF102988.1 /DEF=Homo sapiens Ca ²⁺ -independent phospholipase A2 short isoform (iPLA2) mRNA, complete cds. /FEA=mRNA /GEN=iPLA2 /PROD=Ca ²⁺ -independent phospholipase A2 short isoform /DB_XREF=gi:5305593 /UG=Hs.120360 phospholipase A2, group VI (cytosolic, calcium-independent) /FL=gb:AF102988.1	AF102988		NP_003551

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210648_x_at	1.26E-04	gb:AB047360.1 /DEF=Homo sapiens SNX 3A mRNA for sorting nexin 3A, complete cds. /FEA=mRNA /GEN=SNX 3A /PROD=sorting nexin 3A /DB_XREF=gi:12957159 /UG=Hs.12102 sorting nexin 3 /FL=gb:AB047360.1	AB047360		NP_690041
210667_s_at	6.93E-04	gb:D86062.1 /DEF=Human mRNA for KNP-Ib, complete cds. /FEA=mRNA /GEN=KNP-I /PROD=KNP-Ib /DB_XREF=gi:1545814 /UG=Hs.182423 ES1 (zebrafish) protein, human homolog of /FL=gb:D86062.1	D86062		NP_004640
210679_x_at	0.003171	gb:BC002629.1 /DEF=Homo sapiens, clone MGC:3878, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3878) /DB_XREF=gi:12803588 /UG=Hs.211563 B-cell CLLlymphoma 7A /FL=gb:BC002629.1	BC002629		
210685_s_at	0.004026	gb:AB028839.1 /DEF=Homo sapiens HDNB1(homzygously deleted in neuroblastoma-1)UFD2 mRNA, complete cds. /FEA=mRNA /GEN=HDNB1UFD2 /PROD=homzygously deleted in neuroblastoma-1UFD2 /DB_XREF=gi:13516466 /UG=Hs.24594 ubiquitination factor E4B (homologous to yeast UFD2) /FL=gb:AB028839.1	AB028839		NP_006039
210686_x_at	1.26E-04	gb:BC001407.1 /DEF=Homo sapiens, Similar to cytochrome c-like antigen, clone MGC:2960, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cytochrome c-like antigen /DB_XREF=gi:12655110 /UG=Hs.253070 Homo sapiens, Similar to cytochrome c-like antigen, clone MGC:2960, mRNA, complete cds /FL=gb:BC001407.1	BC001407		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210691_s_at	5.03E-04	gb:AF275803.1 /DEF=Homo sapiens PNAS-107 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-107 /DB_XREF=gi:10834769 /UG=Hs.27258 calcyclin binding protein /FL=gb:AF275803.1	AF275803		NP_055227
210692_s_at	0.001483	gb:BC003163.1 /DEF=Homo sapiens, Similar to selectively expressed in embryonic epithelia protein-1, clone MGC:4220, mRNA, complete cds. /FEA=mRNA /PROD=Similar to selectively expressed in embryonic epithelia protein-1 /DB_XREF=gi:13111980 /UG=Hs.274453 hypothetical protein DKFZp762A227 /FL=gb:BC003163.1	BC003163		
210706_s_at	6.68E-04	gb:BC000213.1 /DEF=Homo sapiens, ring finger protein 24, clone MGC:1815, mRNA, complete cds. /FEA=mRNA /PROD=ring finger protein 24 /DB_XREF=gi:12802985 /UG=Hs.30524 ring finger protein 24 /FL=gb:BC000213.1	BC000213		NP_009150
210715_s_at	0.016934	gb:AF027205.1 /DEF=Homo sapiens Kunitz-type protease inhibitor (kop) mRNA, complete cds. /FEA=mRNA /GEN=kop /PROD=Kunitz-type protease inhibitor /DB_XREF=gi:2598967 /UG=Hs.31439 serine protease inhibitor, Kunitz type, 2 /FL=gb:AF027205.1	AF027205		NP_066925
210716_s_at	1.26E-04	gb:M97501.1 /DEF=Human cytoplasmic linker protein-170 alpha-2 (CLIP-170) mRNA, complete cds. /FEA=mRNA /PROD=cytoplasmic linker protein-170 alpha-2 /DB_XREF=gi:180621 /UG=Hs.31638 restin (Reed-Steinberg cell-expressed intermediate filament-associated protein) /FL=gb:M97501.1	M97501		NP_002947

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210718_s_at	0.037567	gb:AF119889.1 /DEF=Homo sapiens PRO2667 mRNA, complete cds. /FEA=mRNA /PROD=PRO2667 /DB_XREF=gi:7770214 /UG=Hs.321170 Homo sapiens PRO2667 mRNA, complete cds /FL=gb:AF119889.1	AF119889		NP_005812
210756_s_at	0.005768	gb:AF308601.1 /DEF=Homo sapiens NOTCH 2 (N2) mRNA, complete cds. /FEA=mRNA /GEN=N2 /PROD=NOTCH 2 /DB_XREF=gi:11275977 /UG=Hs.8121 Notch (Drosophila) homolog 2 /FL=gb:AF308601.1	AF308601		
210759_s_at	0.033381	gb:M64992.1 /DEF=Human prosomal protein P30-33K (pros-30) mRNA, complete cds. /FEA=mRNA /GEN=pros-30 /PROD=prosomal protein P30-33K /DB_XREF=gi:190446 /UG=Hs.82159 proteasome (prosome, macropain) subunit, alpha type, 1 /FL=gb:M64992.1	M64992		NP_683877
210772_at	0.013223	gb:M88107.1 /DEF=Human formyl peptide receptor (FPR2) mRNA, complete cds. /FEA=mRNA /GEN=FPR2 /PROD=formyl peptide receptor /DB_XREF=gi:189862 /UG=Hs.99855 formyl peptide receptor-like 1 /FL=gb:M76672.1 gb:M84562.1 gb:M88107.1 gb:U81501.1 gb:AF054013.1 gb:NM_001462.1	M88107		NP_001453
210775_x_at	6.79E-04	gb:AB015653.1 /DEF=Homo sapiens mRNA for caspase-9 beta, complete cds. /FEA=mRNA /PROD=caspase-9 beta /DB_XREF=gi:5103139 /UG=Hs.100641 caspase 9, apoptosis-related cysteine protease /FL=gb:AF093130.1 gb:AF110376.1 gb:AB015653.1 gb:AB020979.1	AB015653		NP_127463

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210788_s_at	1.26E-04	gb:AF126782.1 /DEF=Homo sapiens retinal short-chain dehydrogenasereductase retSDR4 mRNA, complete cds. /FEA=mRNA /PROD=retinal short-chain dehydrogenasereductaseretSDR4 /DB_XREF=gi:6318547 /UG=Hs.109201 CGI-86 protein /FL=gb:AF126782.1	AF126782		
210789_x_at	0.042536	gb:L00692.1 /DEF=Human carcinoembryonic antigen (CGM1) mRNA, complete cds. /FEA=mRNA /GEN=CGM1 /PROD=carcinoembryonic antigen /DB_XREF=gi:180226 /UG=Hs.11 carcinoembryonic antigen-related cell adhesion molecule 3 /FL=gb:L00692.1	L00692		NP_001806
210793_s_at	5.03E-04	gb:U41815.1 /DEF=Human nucleoporin 98 (NUP98) mRNA, complete cds. /FEA=mRNA /GEN=NUP98 /PROD=nucleoporin 98 /DB_XREF=gi:1184172 /UG=Hs.112255 nucleoporin 98kD /FL=gb:U41815.1	U41815		NP_624358
210835_s_at	0.016934	gb:AF222711.1 /DEF=Homo sapiens ribeye mRNA, complete cds. /FEA=mRNA /PROD=ribeye /DB_XREF=gi:12034652 /UG=Hs.171391 C-terminal binding protein 2 /FL=gb:AF222711.1 gb:NM_022802.1	AF222711		NP_073713
210840_s_at	0.007779	gb:D29640.1 /DEF=Human mRNA for KIAA0051 gene, complete cds. /FEA=mRNA /GEN=KIAA0051 /DB_XREF=gi:473930 /UG=Hs.1742 IQ motif containing GTPase activating protein 1 /FL=gb:D29640.1	D29640		NP_003861

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210844_x_at	5.03E-04	gb:D14705.1 /DEF=Human mRNA for alpha-catenin, complete cds. /FEA=mRNA /PROD=human alpha-catenin /DB_XREF=gi:415305 /UG=Hs.178452 catenin (cadherin-associated protein), alpha 1 (102kD) /FL=gb:BC000385.1 gb:D14705.1 gb:D13866.1	D14705		NP_001894
210845_s_at	0.00161	gb:U08839.1 /DEF=Human urokinase-type plasminogen activator receptor mRNA, complete cds. /FEA=mRNA /PROD=urokinase-type plasminogen activator receptor /DB_XREF=gi:517197 /UG=Hs.179657 plasminogen activator, urokinase receptor /FL=gb:U08839.1	U08839		NP_002650
210858_x_at	0.013223	gb:U26455.1 /DEF=Human phosphatidylinositol 3-kinase homolog (ATM) mRNA, complete cds. /FEA=mRNA /GEN=ATM /PROD=phosphatidylinositol 3-kinase homolog /DB_XREF=gi:870785 /UG=Hs.194382 ataxia telangiectasia mutated (includes complementation groups A, C and D) /FL=gb:U26455.1	U26455		NP_612150
210859_x_at	0.001618	gb:AF077973.1 /DEF=Homo sapiens clone 36 CLN3 protein (CLN3) mRNA, complete cds. /FEA=mRNA /GEN=CLN3 /PROD=CLN3 protein /DB_XREF=gi:5801864 /UG=Hs.194660 ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease) /FL=gb:AF077973.1	AF077973		NP_000077

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210872_x_at	0.022752	gb:BC001152.1 /DEF=Homo sapiens, growth arrest-specific 7, clone MGC:1347, mRNA, complete cds. /FEA=mRNA /PROD=growth arrest-specific 7 /DB_XREF=gi:12654630 /UG=Hs.226133 growth arrest-specific 7 /FL=gb:BC001152.1	BC001152		NP_005881
210875_s_at	0.001848	gb:U12170.1 /DEF=Human zinc finger homeodomain protein mRNA, complete cds. /FEA=mRNA /PROD=zinc finger homeodomain protein /DB_XREF=gi:529172 /UG=Hs.232068 transcription factor 8 (represses interleukin 2 expression) /FL=gb:U12170.1	U12170		NP_110378
210878_s_at	5.03E-04	gb:BC001202.1 /DEF=Homo sapiens, clone MGC:3185, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3185) /DB_XREF=gi:12654720 /UG=Hs.24125 putative zinc finger protein /FL=gb:BC001202.1	BC001202		NP_057688
210879_s_at	0.002418	gb:AF334812.1 /DEF=Homo sapiens Rab11 interacting protein Rip11a mRNA, complete cds. /FEA=mRNA /PROD=Rab11 interacting protein Rip11a /DB_XREF=gi:13377896 /UG=Hs.24557 KIAA0857 protein /FL=gb:AF334812.1	AF334812		NP_056285
210891_s_at	3.26E-04	gb:AF035737.1 /DEF=Homo sapiens general transcription factor 2-I (GTF2I) mRNA, complete cds. /FEA=mRNA /GEN=GTF2I /PROD=general transcription factor 2-I; alternativesplice product /DB_XREF=gi:2827179 /UG=Hs.278589 general transcription factor II, i /FL=gb:AF035737.1	AF035737		NP_127496

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210907_s_at	1.26E-04	gb:BC002506.1 /DEF=Homo sapiens, programmed cell death 10, clone MGC:1212, mRNA, complete cds. /FEA=mRNA /PROD=programmed cell death 10 /DB_XREF=gi:12803370 /UG=Hs.28866 programmed cell death 10 /FL=gb:BC002506.1	BC002506		NP_665859
210915_x_at	7.58E-04	gb:M15564.1 /DEF=Human T-cell receptor rearranged beta-chain V-region (V-D-J) mRNA, complete cds. /FEA=mRNA /GEN=TCRB /DB_XREF=gi:339011 /UG=Hs.303157 T cell receptor beta locus /FL=gb:M15564.1	M15564		
210926_at	0.00228	gb:AY014272.1 /DEF=Homo sapiens FKSG30 (FKSG30) mRNA, complete cds. /FEA=mRNA /GEN=FKSG30 /PROD=FKSG30 /DB_XREF=gi:12408251 /UG=Hs.315492 Homo sapiens FKSG30 (FKSG30) mRNA, complete cds /FL=gb:AY014272.1	AY014272		
210927_x_at	7.58E-04	gb:BC004239.1 /DEF=Homo sapiens, jumping translocation breakpoint, clone MGC:10274, mRNA, complete cds. /FEA=mRNA /PROD=jumping translocation breakpoint /DB_XREF=gi:13278986 /UG=Hs.323093 Homo sapiens, jumping translocation breakpoint, clone MGC:10274, mRNA, complete cds /FL=gb:BC004239.1	BC004239		NP_006685
210942_s_at	0.010205	gb:AB022918.1 /DEF=Homo sapiens mRNA for alpha2,3-sialyltransferase ST3Gal VI, complete cds. /FEA=mRNA /GEN=ST3Gal VI /PROD=alpha2,3-sialyltransferase ST3Gal VI /DB_XREF=gi:4827246 /UG=Hs.34578 alpha2,3-sialyltransferase /FL=gb:AB022918.1	AB022918		NP_006091

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210943_s_at	1.26E-04	gb:U84744.1 /DEF=Human Chediak-Higashi syndrome protein short isoform (LYST) mRNA, complete cds. /FEA=mRNA /GEN=LYST /PROD=Chediak-Higashi syndrome protein short isoform /DB_XREF=gi:2654473 /UG=Hs.36508 Chediak-Higashi syndrome 1 /FL=gb:U84744.1	U84744		NP_000072
210944_s_at	0.005852	gb:BC003169.1 /DEF=Homo sapiens, Similar to calpain 3, (p94), clone MGC:4403, mRNA, complete cds. /FEA=mRNA /PROD=Similar to calpain 3, (p94) /DB_XREF=gi:13111992 /UG=Hs.40300 calpain 3, (p94) /FL=gb:BC003169.1	BC003169		NP_775113
210950_s_at	1.26E-04	gb:BC003573.1 /DEF=Homo sapiens, farnesyl-diphosphate farnesyltransferase 1, clone MGC:2200, mRNA, complete cds. /FEA=mRNA /PROD=farnesyl-diphosphate farnesyltransferase 1 /DB_XREF=gi:13097746 /UG=Hs.48876 farnesyl-diphosphate farnesyltransferase 1 /FL=gb:BC003573.1	BC003573		NP_004453
210951_x_at	3.26E-04	gb:AF125393.1 /DEF=Homo sapiens Rab27 isoform mRNA, complete cds. /FEA=mRNA /PROD=Rab27 isoform /DB_XREF=gi:5410356 /UG=Hs.50477 RAB27A, member RAS oncogene family /FL=gb:AF125393.1	AF125393		NP_004571

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210965_x_at	0.002418	gb:BC001274.1 /DEF=Homo sapiens, Similar to cell division cycle 2-like 5 (cholinesterase-related cell division controller), clone MGC:5091, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cell division cycle 2-like 5(cholinesterase-related cell division controller) /DB_XREF=gi:12654860 /UG=Hs.59498 cell division cycle 2-like 5 (cholinesterase-related cell division controller) /FL=gb:BC001274.1	BC001274		NP_112557
210966_x_at	0.00545	gb:BC001460.1 /DEF=Homo sapiens, Similar to hypothetical protein FLJ10378, clone MGC:2954, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein FLJ10378 /DB_XREF=gi:12655204 /UG=Hs.6214 KIAA0731 protein /FL=gb:BC001460.1	BC001460		NP_056130
210968_s_at	0.02145	gb:AF333336.1 /DEF=Homo sapiens testis specific reticulon 5 protein mRNA, complete cds. /FEA=mRNA /PROD=testis specific reticulon 5 protein /DB_XREF=gi:13377627 /UG=Hs.65450 reticulon 4 /FL=gb:AF333336.1	AF333336		NP_722550
210969_at	0.003529	gb:AF118089.1 /DEF=Homo sapiens PRO2042 mRNA, complete cds. /FEA=mRNA /PROD=PRO2042 /DB_XREF=gi:6650823 /UG=Hs.69171 protein kinase C-like 2 /FL=gb:AF118089.1	AF118089		NP_006247
210972_x_at	2.05E-04	gb:M15565.1 /DEF=Human T-cell receptor rearranged alpha-chain V-region (V-D-J) mRNA, complete cds. /FEA=mRNA /GEN=TCRA /DB_XREF=gi:338765 /UG=Hs.74647 Human T-cell receptor active alpha-chain mRNA from JM cell line, complete cds /FL=gb:M15565.1	M15565		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
210980_s_at	1.26E-04	gb:U47674.1 /DEF=Homo sapiens putative heart protein PHP mRNA, complete cds. /FEA=mRNA /PROD=PHP /DB_XREF=gi:3860239 /UG=Hs.75811 N-acylsphingosine amidohydrolase (acid ceramidase) /FL=gb:U47674.1	U47674		NP_808592
210981_s_at	0.001618	gb:AF040751.1 /DEF=Homo sapiens G protein-coupled receptor kinase 6, splice variant B (GRK6) mRNA, complete cds. /FEA=mRNA /GEN=GRK6 /PROD=G protein-coupled receptor kinase 6, splicevariant B /DB_XREF=gi:3005015 /UG=Hs.76297 G protein-coupled receptor kinase 6 /FL=gb:AF040751.1	AF040751		NP_002073
210986_s_at	0.00161	gb:Z24727.1 /DEF=H.sapiens tropomyosin isoform mRNA, complete CDS. /FEA=mRNA /PROD=tropomyosin isoform /DB_XREF=gi:854188 /UG=Hs.77899 tropomyosin 1 (alpha) /FL=gb:Z24727.1	Z24727		NP_000357
210987_x_at	1.26E-04	gb:M19267.1 /DEF=Human tropomyosin mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:339943 /UG=Hs.77899 tropomyosin 1 (alpha) /FL=gb:M19267.1	M19267		
210996_s_at	3.26E-04	gb:U43430.1 /DEF=Human epsilon isoform 14-3-3 protein mRNA, complete cds. /FEA=mRNA /PROD=14-3-3 protein /DB_XREF=gi:4096984 /UG=Hs.79474 tyrosine 3-monooxygenasetryptophan 5-monooxygenase activation protein, epsilon polypeptide /FL=gb:U43399.1 gb:U43430.1	U43430		NP_006752

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211015_s_at	2.05E-04	gb:L12723.1 /DEF=Human heat shock protein 70 (hsp70) mRNA, complete cds. /FEA=mRNA /GEN=hsp70 /PROD=heat shock protein 70 /DB_XREF=gi:292159 /UG=Hs.90093 heat shock 70kD protein 4 /FL=gb:L12723.1	L12723		
211025_x_at	1.26E-04	gb:BC006229.1 /DEF=Homo sapiens, cytochrome c oxidase subunit Vb, clone MGC:10622, mRNA, complete cds. /FEA=mRNA /PROD=cytochrome c oxidase subunit Vb /DB_XREF=gi:13623258 /FL=gb:BC006229.1	BC006229		NP_001853
211026_s_at	0.004341	gb:BC006230.1 /DEF=Homo sapiens, lysophospholipase-like, clone MGC:10338, mRNA, complete cds. /FEA=mRNA /PROD=lysophospholipase-like /DB_XREF=gi:13623260 /FL=gb:BC006230.1	BC006230		NP_009214
211034_s_at	0.005339	gb:BC006270.1 /DEF=Homo sapiens, clone MGC:11291, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:11291) /DB_XREF=gi:13623331 /FL=gb:BC006270.1	BC006270		
211040_x_at	0.004341	gb:BC006325.1 /DEF=Homo sapiens, G-2 and S-phase expressed 1, clone MGC:12560, mRNA, complete cds. /FEA=mRNA /PROD=G-2 and S-phase expressed 1 /DB_XREF=gi:13623450 /FL=gb:BC006325.1	BC006325		NP_057510
211043_s_at	0.001116	gb:BC006332.1 /DEF=Homo sapiens, clathrin, light polypeptide (Lcb), clone MGC:12930, mRNA, complete cds. /FEA=mRNA /PROD=clathrin, light polypeptide (Lcb) /DB_XREF=gi:13623458 /FL=gb:BC006332.1	BC006332		NP_009028

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211047_x_at	0.013223	gb:BC006337.1 /DEF=Homo sapiens, clone MGC:12798, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:12798) /DB_XREF=gi:13623468 /FL=gb:BC006337.1	BC006337		NP_067586
211058_x_at	5.03E-04	gb:BC006379.1 /DEF=Homo sapiens, tubulin alpha 1, clone MGC:12832, mRNA, complete cds. /FEA=mRNA /PROD=tubulin alpha 1 /DB_XREF=gi:13623540 /FL=gb:BC006379.1	BC006379		NP_006073
211060_x_at	0.002418	gb:BC006383.1 /DEF=Homo sapiens, clone MGC:12663, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:12663) /DB_XREF=gi:13623546 /FL=gb:BC006383.1	BC006383		NP_003792
211067_s_at	0.016074	gb:BC006454.1 /DEF=Homo sapiens, growth arrest-specific 7, clone MGC:1348, mRNA, complete cds. /FEA=mRNA /PROD=growth arrest-specific 7 /DB_XREF=gi:13623656 /FL=gb:BC006454.1	BC006454		NP_005881
211068_x_at	0.010205	gb:BC006456.1 /DEF=Homo sapiens, clone MGC:1426, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:1426) /DB_XREF=gi:13623660 /FL=gb:BC006456.1	BC006456		
211072_x_at	5.03E-04	gb:BC006481.1 /DEF=Homo sapiens, tubulin alpha 1, clone MGC:4387, mRNA, complete cds. /FEA=mRNA /PROD=tubulin alpha 1 /DB_XREF=gi:13623706 /FL=gb:BC006481.1	BC006481		NP_006073
211073_x_at	0.041055	gb:BC006483.1 /DEF=Homo sapiens, ribosomal protein L3, clone MGC:4393, mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein L3 /DB_XREF=gi:13623708 /FL=gb:BC006483.1	BC006483		NP_000958

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211074_at	0.003171	gb:AF000381.1 /DEF=Homo sapiens non-functional folate binding protein mRNA, complete cds. /FEA=mRNA /PROD=non-functional folate binding protein /DB_XREF=gi:2565195 /FL=gb:AF000381.1	AF000381		
211075_s_at	1.26E-04	gb:Z25521.1 /DEF=H.sapiens integrin associated protein mRNA, complete CDS,. /FEA=mRNA /PROD=integrin associated protein /DB_XREF=gi:396704 /FL=gb:Z25521.1	Z25521		NP_001768
211085_s_at	5.03E-04	gb:Z25430.1 /DEF=H.sapiens protein-serinethreonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serinethreonine kinase /DB_XREF=gi:405738 /FL=gb:Z25430.1	Z25430		NP_006273
211087_x_at	0.011725	gb:Z25432.1 /DEF=H.sapiens protein-serinethreonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serinethreonine kinase /DB_XREF=gi:405742 /FL=gb:Z25432.1	Z25432		NP_620583
211097_s_at	0.009689	gb:BC003111.1 /DEF=Homo sapiens, Similar to pre-B-cell leukemia transcription factor 2, clone MGC:2174, mRNA, complete cds. /FEA=mRNA /PROD=Similar to pre-B-cell leukemia transcriptionfactor 2 /DB_XREF=gi:13111886 /UG=Hs.93728 pre-B-cell leukemia transcription factor 2 /FL=gb:BC003111.1	BC003111		
211098_x_at	6.68E-04	gb:AF277194.1 /DEF=Homo sapiens PNAS-136 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-136 /DB_XREF=gi:12751112 /UG=Hs.93832 putative membrane protein /FL=gb:AF277194.1	AF277194		NP_061899

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211100_x_at	0.026891	gb:U82278.1 /DEF=Human immunoglobulin-like transcript 1c mRNA, complete cds. /FEA=mRNA /PROD=immunoglobulin-like transcript 1c /DB_XREF=gi:1907322 /UG=Hs.94498 leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 /FL=gb:U82278.1 gb:AF025531.1	U82278		NP_006857
211121_s_at	0.002989	gb:AF180527.1 /DEF=Homo sapiens p22Dokdel (DOKDEL) mRNA, complete cds. /FEA=mRNA /GEN=DOKDEL /PROD=p22Dokdel /DB_XREF=gi:6606314 /UG=Hs.103854 docking protein 1, 62kD (downstream of tyrosine kinase 1) /FL=gb:AF180527.1	AF180527		NP_001372
211137_s_at	0.0375	gb:AF189723.2 /DEF=Homo sapiens calcium transport ATPase ATP2C1 (ATP2C1A) mRNA, complete cds. /FEA=mRNA /GEN=ATP2C1A /PROD=calcium transport ATPase ATP2C1 /DB_XREF=gi:6826913 /UG=Hs.106778 ATPase, Ca++ transporting, type 2C, member 1 /FL=gb:AF189723.2	AF189723		NP_055197
211144_x_at	0.026891	gb:M30894.1 /DEF=Human T-cell receptor Ti rearranged gamma-chain mRNA V-J-C region, complete cds. /FEA=mRNA /GEN=CD3G /DB_XREF=gi:339406 /UG=Hs.112259 T cell receptor gamma locus /FL=gb:M30894.1	M30894		
211160_x_at	0.033491	gb:M95178.1 /DEF=Human non-muscle alpha-actinin mRNA, complete cds. /FEA=mRNA /GEN=ACTN1 /PROD=alpha-actinin /DB_XREF=gi:178051 /UG=Hs.119000 actinin, alpha 1 /FL=gb:M95178.1	M95178		NP_001093

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211202_s_at	0.013223	gb:AF087481.1 /DEF=Homo sapiens retinoblastoma binding protein 2 homolog 1 (RBBP2H1) mRNA, complete cds. /FEA=mRNA /GEN=RBBP2H1 /PROD=retinoblastoma binding protein 2 homolog 1 /DB_XREF=gi:4322487 /UG=Hs.143323 putative DNAchromatin binding motif /FL=gb:AF087481.1	AF087481		NP_006609
211220_s_at	0.038535	gb:BC005329.1 /DEF=Homo sapiens, Similar to heat shock transcription factor 2, clone MGC:12415, mRNA, complete cds. /FEA=mRNA /PROD=Similar to heat shock transcription factor 2 /DB_XREF=gi:13529106 /UG=Hs.158195 heat shock transcription factor 2 /FL=gb:BC005329.1	BC005329		NP_004497
211251_x_at	0.007052	gb:U78774.1 /DEF=Human NFY-C mRNA, complete cds. /FEA=mRNA /PROD=NFY-C /DB_XREF=gi:2327008 /UG=Hs.168157 nuclear transcription factor Y, gamma /FL=gb:U78774.1	U78774		NP_055038
211252_x_at	0.013575	gb:U36759.1 /DEF=Human pre-T cell receptor alpha-type chain precursor, mRNA, complete cds. /FEA=mRNA /PROD=pre-T cell receptor alpha-type chain precursor /DB_XREF=gi:1127580 /UG=Hs.169002 Human pre TCR alpha mRNA, partial cds /FL=gb:U36759.1	U36759		NP_612153
211255_x_at	0.004541	gb:AF064605.1 /DEF=Homo sapiens KE05 protein mRNA, complete cds. /FEA=mRNA /PROD=KE05 protein /DB_XREF=gi:3152663 /UG=Hs.169681 death effector domain-containing /FL=gb:AF064605.1	AF064605		NP_127491

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211275_s_at	1.26E-04	gb:AF087942.1 /DEF=Homo sapiens glycogenin-1L mRNA, complete cds. /FEA=mRNA /PROD=glycogenin-1L /DB_XREF=gi:5814084 /UG=Hs.174071 glycogenin /FL=gb:AF087942.1	AF087942		NP_004121
211284_s_at	0.041055	gb:BC000324.1 /DEF=Homo sapiens, Similar to granulin, clone MGC:8480, mRNA, complete cds. /FEA=mRNA /PROD=Similar to granulin /DB_XREF=gi:12653114 /UG=Hs.180577 granulin /FL=gb:BC000324.1	BC000324		NP_002078
211285_s_at	0.041055	gb:U84404.1 /DEF=Human E6-associated protein E6-APubiquitin-protein ligase (UBE3A) mRNA, alternatively spliced, complete cds. /FEA=mRNA /GEN=UBE3A /PROD=E6-associated protein E6-APubiquitin-proteinligase /DB_XREF=gi:1872513 /UG=Hs.180686 ubiquitin protein ligase E3A (human papilloma virus E6-associated protein, Angelman syndrome) /FL=gb:AF273050.1 gb:U84404.1	U84404		NP_570854
211286_x_at	0.009716	gb:L29349.1 /DEF=Homo sapiens granulocyte-macrophage colony-stimulating factor receptor alpha-subunit 3 (GM-CSF-RA3) mRNA, complete cds. /FEA=mRNA /GEN=GM-CSF-RA3 /PROD=granulocyte-macrophage colony-stimulating factorreceptor alpha-subunit 3 /DB_XREF=gi:460284 /UG=Hs.182378 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage) /FL=gb:L29349.1	L29349		NP_758452

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211287_x_at	1.26E-04	gb:M64445.1 /DEF=Human GM-CSF receptor mRNA, complete cds. /FEA=mRNA /GEN=GM-CSF receptor /PROD=GM-CSF receptor /DB_XREF=gi:183361 /UG=Hs.182378 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage) /FL=gb:M64445.1	M64445		NP_758452
211296_x_at	0.026891	gb:AB009010.1 /DEF=Homo sapiens mRNA for polyubiquitin UbC, complete cds. /FEA=mRNA /GEN=UbC1 /PROD=polyubiquitin UbC /DB_XREF=gi:2647407 /UG=Hs.183704 ubiquitin C /FL=gb:BC000449.1 gb:AB009010.1	AB009010		NP_066289
211297_s_at	1.28E-04	gb:L20320.1 /DEF=Human protein serinethreonine kinase stk1 mRNA, complete cds. /FEA=mRNA /PROD=protein serinethreonine kinase /DB_XREF=gi:348242 /UG=Hs.184298 cyclin-dependent kinase 7 (homolog of Xenopus MO15 cdk-activating kinase) /FL=gb:L20320.1	L20320		NP_001790
211307_s_at	0.005276	gb:U43677.1 /DEF=Human Fc alpha receptor, splice variant FcalphaR a.3 (CD89) mRNA, complete cds. /FEA=mRNA /GEN=CD89 /PROD=Fc alpha receptor /DB_XREF=gi:1439557 /UG=Hs.193122 Fc fragment of IgA, receptor for /FL=gb:U43677.1 gb:D87854.1 gb:D87856.1	U43677		NP_579814
211318_s_at	2.05E-04	gb:U85943.1 /DEF=Homo sapiens mRNA-associated protein mrnp41 mRNA, complete cds. /FEA=mRNA /PROD=mRNA-associated protein mrnp41 /DB_XREF=gi:2231591 /UG=Hs.196209 RAE1 (RNA export 1, S.pombe) homolog /FL=gb:U85943.1	U85943		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211325_x_at	0.007779	gb:U72518.1 /DEF=Human destrin-2 pseudogene mRNA, complete cds. /FEA=mRNA /PROD=destrin-2 /DB_XREF=gi:1673523 /UG=Hs.199299 Human destrin-2 pseudogene mRNA, complete cds /FL=gb:U72518.1	U72518		
211342_x_at	0.02145	gb:BC004354.1 /DEF=Homo sapiens, trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit), clone MGC:3120, mRNA, complete cds. /FEA=mRNA /PROD=trinucleotide repeat containing 11(THR-associated protein, 230 kDa subunit) /DB_XREF=gi:13279313 /UG=Hs.211607 trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit) /FL=gb:BC004354.1	BC004354		
211355_x_at	0.013274	gb:U52914.1 /DEF=Human B219OB receptor isoform HuB219.3 precursor mRNA, complete cds. /FEA=mRNA /PROD=B219OB receptor isoform HuB219.3 precursor /DB_XREF=gi:1279904 /UG=Hs.226627 leptin receptor /FL=gb:U52914.1	U52914		NP_002294
211383_s_at	2.05E-04	gb:AL136827.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434F2427 (from clone DKFZp434F2427); complete cds. /FEA=mRNA /GEN=DKFZp434F2427 /PROD=hypothetical protein /DB_XREF=gi:6807664 /UG=Hs.27207 KIAA0982 protein /FL=gb:AL136827.1	AL136827		NP_054742
211404_s_at	1.26E-04	gb:BC004371.1 /DEF=Homo sapiens, clone MGC:10449, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10449) /DB_XREF=gi:13325115 /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2 /FL=gb:BC004371.1	BC004371		NP_001633

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211416_x_at	0.033381	gb:L20492.1 /DEF=Human gamma-glutamyl transpeptidase mRNA, complete cds. /FEA=mRNA /PROD=gamma-glutamyl transpeptidase /DB_XREF=gi:306752 /UG=Hs.284380 gamma-glutamyltransferase 1 /FL=gb:L20492.1	L20492		NP_842564
211417_x_at	1.26E-04	gb:L20493.1 /DEF=Human gamma-glutamyl transpeptidase mRNA, complete cds. /FEA=mRNA /PROD=gamma-glutamyl transpeptidase /DB_XREF=gi:306754 /UG=Hs.284380 gamma-glutamyltransferase 1 /FL=gb:L20493.1	L20493		
211426_x_at	0.011725	gb:U40038.1 /DEF=Human GTP-binding protein alpha q subunit (GNAQ) mRNA, complete cds. /FEA=mRNA /GEN=GNAQ /PROD=GTP-binding protein alpha q subunit /DB_XREF=gi:1181670 /UG=Hs.296261 guanine nucleotide binding protein (G protein), q polypeptide /FL=gb:U40038.1	U40038		NP_002063
211429_s_at	0.041055	gb:AF119873.1 /DEF=Homo sapiens PRO2275 mRNA, complete cds. /FEA=mRNA /PROD=PRO2275 /DB_XREF=gi:7770182 /UG=Hs.297681 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1 /FL=gb:AF119873.1	AF119873		NP_000286
211433_x_at	0.00161	gb:AL583909.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761J197 (from clone DKFZp761J197); complete cds. /FEA=mRNA /GEN=DKFZp761J197 /PROD=hypothetical protein /DB_XREF=gi:13093772 /UG=Hs.301696 hypothetical protein FLJ11560 /FL=gb:AL583909.1	AL583909		NP_079458

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211454_x_at	5.03E-04	gb:AF336878.1 /DEF=Homo sapiens FKSG51 (FKSG51) mRNA, complete cds. /FEA=mRNA /GEN=FKSG51 /PROD=FKSG51 /DB_XREF=gi:13384184 /UG=Hs.326752 Homo sapiens FKSG51 (FKSG51) mRNA, complete cds /FL=gb:AF336878.1	AF336878		
211471_s_at	0.028067	gb:AF133588.1 /DEF=Homo sapiens RAB36 (RAB36) mRNA, complete cds. /FEA=mRNA /GEN=RAB36 /PROD=RAB36 /DB_XREF=gi:6049163 /UG=Hs.38772 RAB36, member RAS oncogene family /FL=gb:AF133588.1	AF133588		NP_004905
211474_s_at	0.005852	gb:BC004948.1 /DEF=Homo sapiens, clone MGC:10846, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10846) /DB_XREF=gi:13436313 /UG=Hs.41072 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 6 /FL=gb:BC004948.1	BC004948		
211475_s_at	1.26E-04	gb:AF116273.1 /DEF=Homo sapiens Bag1 protein variant mRNA, complete cds. /FEA=mRNA /PROD=Bag1 protein variant /DB_XREF=gi:4583370 /UG=Hs.41714 BCL2-associated athanogene /FL=gb:AF116273.1	AF116273		NP_004314
211505_s_at	1.26E-04	gb:AL136601.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564J1516 (from clone DKFZp564J1516); complete cds. /FEA=mRNA /GEN=DKFZp564J1516 /PROD=hypothetical protein /DB_XREF=gi:13276702 /UG=Hs.6113 stau6n (Drosophila, RNA-binding protein) /FL=gb:AL136601.1	AL136601		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211509_s_at	1.26E-04	gb:AB015639.1 /DEF=Homo sapiens ASY mRNA, complete cds. /FEA=mRNA /GEN=ASY /DB_XREF=gi:5821139 /UG=Hs.65450 reticulon 4 /FL=gb:AB015639.1	AB015639		NP_722550
211529_x_at	0.005852	gb:M90684.1 /DEF=Human lymphocyte antigen (HLA-G2.1) mRNA, complete cds. /FEA=mRNA /GEN=HLA-G /PROD=b2 microglobulin /DB_XREF=gi:188467 /UG=Hs.73885 HLA-G histocompatibility antigen, class I, G /FL=gb:M90684.1	M90684		NP_002118
211542_x_at	0.041055	gb:BC004334.1 /DEF=Homo sapiens, ribosomal protein S10, clone MGC:10943, mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein S10 /DB_XREF=gi:13279259 /UG=Hs.76230 ribosomal protein S10 /FL=gb:BC004334.1	BC004334		NP_001005
211546_x_at	1.36E-04	gb:L36674.1 /DEF=Human (clone 2-5) synuclein (NACP) mRNA, complete cds. /FEA=mRNA /GEN=NACP /PROD=synuclein /DB_XREF=gi:556211 /UG=Hs.76930 synuclein, alpha (non A4 component of amyloid precursor) /FL=gb:L36674.1	L36674		NP_009292
211548_s_at	0.045541	gb:J05594.1 /DEF=Homo sapiens NAD+-dependent 15-hydroxyprostaglandin dehydrogenase (PDGH) mRNA, complete cds. /FEA=mRNA /GEN=PGDH /PROD=NAD+-dependent 15-hydroxyprostaglandin dehydrogenase /DB_XREF=gi:1203981 /UG=Hs.77348 hydroxyprostaglandin dehydrogenase 15-(NAD) /FL=gb:J05594.1	J05594		NP_000851

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211569_s_at	0.011725	gb:AF001903.1 /DEF=Human 3-hydroxyacyl-CoA dehydrogenase, isoform 2 mRNA, complete cds. /FEA=mRNA /PROD=3-hydroxyacyl-CoA dehydrogenase, isoform 2 /DB_XREF=gi:2078328 /UG=Hs.8110 L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain /FL=gb:AF001903.1	AF001903		NP_005318
211571_s_at	0.026891	gb:D32039.1 /DEF=Human pgH3 mRNA for proteoglycan PG-M(V3), complete cds. /FEA=mRNA /GEN=pgH3 /PROD=proteoglycan PG-M(V3) /DB_XREF=gi:1008912 /UG=Hs.81800 chondroitin sulfate proteoglycan 2 (versican) /FL=gb:D32039.1	D32039		NP_004376
211574_s_at	0.027418	gb:D84105.1 /DEF=Human CD46 mRNA, complete cds. /FEA=mRNA /PROD=CD46 /DB_XREF=gi:1256700 /UG=Hs.83532 membrane cofactor protein (CD46, trophoblast-lymphocyte cross-reactive antigen) /FL=gb:D84105.1	D84105		NP_758871
211581_x_at	0.007779	gb:AF000426.1 /DEF=Homo sapiens LST1 mRNA, cLST1E splice variant, complete cds. /FEA=mRNA /GEN=LST1 /DB_XREF=gi:2145067 /UG=Hs.88411 lymphocyte antigen 117 /FL=gb:AF000426.1	AF000426		NP_009092
211582_x_at	0.013223	gb:AF000424.1 /DEF=Homo sapiens LST1 mRNA, cLST1C splice variant, complete cds. /FEA=mRNA /GEN=LST1 /DB_XREF=gi:2145063 /UG=Hs.88411 lymphocyte antigen 117 /FL=gb:AF000424.1	AF000424		NP_009092
211609_x_at	0.00161	gb:U51007.1 /DEF=Human 26S protease subunit S5a mRNA, complete cds. /FEA=mRNA /PROD=26S protease subunit S5a /DB_XREF=gi:1256400 /FL=gb:U51007.1	U51007		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211623_s_at	0.005852	gb:M30448.1 /DEF=Human casein kinase II beta subunit mRNA, complete cds. /FEA=mRNA /GEN=CSNK2B /DB_XREF=gi:181154 /FL=gb:M30448.1	M30448		
211654_x_at	0.041055	gb:M17565.1 /DEF=Human MHC class II DQ-beta associated with DRw6, DQw1 protein, complete cds. /FEA=mRNA /GEN=HLA-DQB1 /DB_XREF=gi:188188 /FL=gb:M17565.1	M17565		NP_002114
211657_at	2.05E-04	gb:M18728.1 /DEF=Human nonspecific crossreacting antigen mRNA, complete cds. /FEA=mRNA /GEN=NCA; NCA; NCA /PROD=non-specific cross reacting antigen /DB_XREF=gi:189084 /FL=gb:M18728.1	M18728		NP_002474
211666_x_at	0.005852	gb:L22453.1 /DEF=Homo sapiens HIV-1 TAR RNA binding protein (TARBP-b) mRNA, complete cds. /FEA=mRNA /GEN=TARBP-b /DB_XREF=gi:347963 /FL=gb:L22453.1	L22453		NP_000958
211671_s_at	0.005852	gb:U01351.1 /DEF=Human glucocorticoid receptor alpha-2 mRNA, complete cds. /FEA=mRNA /PROD=glucocorticoid receptor alpha-2 /DB_XREF=gi:458656 /FL=gb:U01351.1	U01351		NP_000167
211676_s_at	1.26E-04	gb:AF056979.1 /DEF=Homo sapiens clone YAN1 interferon-gamma receptor mRNA, complete cds. /FEA=mRNA /PROD=interferon-gamma receptor /DB_XREF=gi:13562048 /FL=gb:AF056979.1	AF056979		NP_000407
211684_s_at	3.26E-04	gb:AF250307.1 /DEF=Homo sapiens cytoplasmic dynein intermediate chain 2C mRNA, complete cds. /FEA=mRNA /PROD=cytoplasmic dynein intermediate chain 2C /DB_XREF=gi:13649464 /FL=gb:AF250307.1	AF250307		NP_001369

Gen Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
211685_s_at	0.007052	gb:AF251061.1 /DEF=Homo sapiens neurocalcin mRNA, complete cds. /FEA=mRNA /PROD=neurocalcin /DB_XREF=gi:13625183 /FL=gb:AF251061.1	AF251061		NP_114430
211693_at	0.009689	gb:AF343666.1 /DEF=Homo sapiens translocation associated fusion protein IRTA1IGA1 (IRTA1IGHA1) mRNA, complete cds. /FEA=mRNA /GEN=IRTA1IGHA1 /PROD=translocation associated fusion protein IRTA1IGA1 /DB_XREF=gi:13591717 /FL=gb:AF343666.1	AF343666		
211696_x_at	0.00228	gb:AF349114.1 /DEF=Homo sapiens beta globin chain variant (HBB) mRNA, complete cds. /FEA=mRNA /GEN=HBB /PROD=beta globin chain variant /DB_XREF=gi:13549111 /FL=gb:AF349114.1	AF349114		NP_000509
211699_x_at	1.28E-04	gb:AF349571.1 /DEF=Homo sapiens hemoglobin alpha-1 globin chain (HBA1) mRNA, complete cds. /FEA=mRNA /GEN=HBA1 /PROD=hemoglobin alpha-1 globin chain /DB_XREF=gi:13650073 /FL=gb:AF349571.1	AF349571		NP_000549
211702_s_at	1.26E-04	gb:AF350251.1 /DEF=Homo sapiens ubiquitin specific protease mRNA, complete cds. /FEA=mRNA /PROD=ubiquitin specific protease /DB_XREF=gi:13560796 /FL=gb:AF350251.1	AF350251		NP_115971
211703_s_at	5.03E-04	gb:AF353990.1 /DEF=Homo sapiens beta-amyloid binding protein precursor (BBP) mRNA, complete cds. /FEA=mRNA /GEN=BBP /PROD=beta-amyloid binding protein precursor /DB_XREF=gi:13625458 /FL=gb:AF353990.1	AF353990		NP_114416

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211710_x_at	0.010205	gb:BC005817.1 /DEF=Homo sapiens, ribosomal protein L4, clone MGC:11073, mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein L4 /DB_XREF=gi:13543303 /FL=gb:BC005817.1	BC005817		NP_000959
211711_s_at	0.013223	gb:BC005821.1 /DEF=Homo sapiens, phosphatase and tensin homolog (mutated in multiple advanced cancers 1), clone MGC:11227, mRNA, complete cds. /FEA=mRNA /PROD=phosphatase and tensin homolog (mutated in multiple advanced cancers 1) /DB_XREF=gi:13543309 /FL=gb:BC005821.1	BC005821		NP_000305
211725_s_at	1.26E-04	gb:BC005884.1 /DEF=Homo sapiens, clone MGC:4736, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:4736) /DB_XREF=gi:13543452 /FL=gb:BC005884.1	BC005884		
211727_s_at	0.02145	gb:BC005895.1 /DEF=Homo sapiens, COX11 (yeast) homolog, cytochrome c oxidase assembly protein, clone MGC:14469, mRNA, complete cds. /FEA=mRNA /PROD=COX11 (yeast) homolog, cytochrome c oxidase assembly protein /DB_XREF=gi:13543474 /FL=gb:BC005895.1	BC005895		NP_004366
211729_x_at	0.003171	gb:BC005902.1 /DEF=Homo sapiens, Similar to biliverdin reductase A, clone MGC:14491, mRNA, complete cds. /FEA=mRNA /PROD=Similar to biliverdin reductase A /DB_XREF=gi:13543488 /FL=gb:BC005902.1	BC005902		NP_000703

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211730_s_at	0.001116	gb:BC005903.1 /DEF=Homo sapiens, polymerase (RNA) II (DNA directed) polypeptide L (7.6kD), clone MGC:14494, mRNA, complete cds. /FEA=mRNA /PROD=polymerase (RNA) II (DNA directed) polypeptide L(7.6kD) /DB_XREF=gi:13543491 /FL=gb:BC005903.1	BC005903		NP_066951
211733_x_at	0.010205	gb:BC005911.1 /DEF=Homo sapiens, sterol carrier protein 2, clone MGC:14505, mRNA, complete cds. /FEA=mRNA /PROD=sterol carrier protein 2 /DB_XREF=gi:13543502 /FL=gb:BC005911.1	BC005911		NP_002970
211741_x_at	0.041795	gb:BC005924.1 /DEF=Homo sapiens, pregnancy specific beta-1-glycoprotein 3, clone MGC:14526, mRNA, complete cds. /FEA=mRNA /PROD=pregnancy specific beta-1-glycoprotein 3 /DB_XREF=gi:13543532 /FL=gb:BC005924.1	BC005924		NP_066296
211742_s_at	0.001116	gb:BC005926.1 /DEF=Homo sapiens, ecotropic viral integration site 2B, clone MGC:14529, mRNA, complete cds. /FEA=mRNA /PROD=ecotropic viral integration site 2B /DB_XREF=gi:13543535 /FL=gb:BC005926.1	BC005926		NP_006486
211744_s_at	1.26E-04	gb:BC005930.1 /DEF=Homo sapiens, Similar to CD58 antigen, (lymphocyte function-associated antigen 3), clone MGC:14538, mRNA, complete cds. /FEA=mRNA /PROD=Similar to CD58 antigen, (lymphocytefunction-associated antigen 3) /DB_XREF=gi:13543544 /FL=gb:BC005930.1	BC005930		NP_001770

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211745_x_at	7.58E-04	gb:BC005931.1 /DEF=Homo sapiens, hemoglobin, alpha 2, clone MGC:14541, mRNA, complete cds. /FEA=mRNA /PROD=hemoglobin, alpha 2 /DB_XREF=gi:13543547 /FL=gb:BC005931.1	BC005931		NP_000508
211746_x_at	0.004341	gb:BC005932.1 /DEF=Homo sapiens, proteasome (prosome, macropain) subunit, alpha type, 1, clone MGC:14542, mRNA, complete cds. /FEA=mRNA /PROD=proteasome (prosome, macropain) subunit, alphas type, 1 /DB_XREF=gi:13543550 /FL=gb:BC005932.1	BC005932		NP_683877
211749_s_at	0.026891	gb:BC005941.1 /DEF=Homo sapiens, Similar to vesicle-associated membrane protein 3, clone MGC:14563, mRNA, complete cds. /FEA=mRNA /PROD=Similar to vesicle-associated membrane protein3 /DB_XREF=gi:13543573 /FL=gb:BC005941.1	BC005941		NP_004772
211755_s_at	0.001116	gb:BC005960.1 /DEF=Homo sapiens, Similar to ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit b, isoform 1, clone MGC:14609, mRNA, complete cds. /FEA=mRNA /PROD=Similar to ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit b, isoform 1 /DB_XREF=gi:13543617 /FL=gb:BC005960.1	BC005960		NP_001679
211758_x_at	0.007779	gb:BC005968.1 /DEF=Homo sapiens, ATP binding protein associated with cell differentiation, clone MGC:14620, mRNA, complete cds. /FEA=mRNA /PROD=ATP binding protein associated with celldifferentiation /DB_XREF=gi:13543638 /FL=gb:BC005968.1	BC005968		NP_005774

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211760_s_at	1.26E-04	gb:BC005974.1 /DEF=Homo sapiens, vesicle-associated membrane protein 4, clone MGC:14658, mRNA, complete cds. /FEA=mRNA /PROD=vesicle-associated membrane protein 4 /DB_XREF=gi:13543647 /FL=gb:BC005974.1	BC005974		NP_003753
211763_s_at	7.58E-04	gb:BC005979.1 /DEF=Homo sapiens, ubiquitin-conjugating enzyme E2B (RAD6 homolog), clone MGC:14670, mRNA, complete cds. /FEA=mRNA /PROD=ubiquitin-conjugating enzyme E2B (RAD6 homolog) /DB_XREF=gi:13543659 /FL=gb:BC005979.1	BC005979		NP_003328
211764_s_at	0.003171	gb:BC005980.1 /DEF=Homo sapiens, ubiquitin-conjugating enzyme E2D 1 (homologous to yeast UBC45), clone MGC:14673, mRNA, complete cds. /FEA=mRNA /PROD=ubiquitin-conjugating enzyme E2D 1 (homologousto yeast UBC45) /DB_XREF=gi:13543662 /FL=gb:BC005980.1	BC005980		NP_003329
211776_s_at	0.00161	gb:BC006141.1 /DEF=Homo sapiens, Similar to erythrocyte protein band 4.1-like 3, clone MGC:13087, mRNA, complete cds. /FEA=mRNA /PROD=Similar to erythrocyte protein band 4.1-like 3 /DB_XREF=gi:13544008 /FL=gb:BC006141.1	BC006141		NP_036439
211779_x_at	1.26E-04	gb:BC006155.1 /DEF=Homo sapiens, clone MGC:13188, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:13188) /DB_XREF=gi:13544040 /FL=gb:BC006155.1	BC006155		NP_036437

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211782_at	0.031213	gb:BC006170.1 /DEF=Homo sapiens, Similar to iduronate 2-sulfatase (Hunter syndrome), clone MGC:13238, mRNA, complete cds. /FEA=mRNA /PROD=Similar to iduronate 2-sulfatase (Huntersyndrome) /DB_XREF=gi:13544076 /FL=gb:BC006170.1	BC006170		NP_006114
211783_s_at	0.001116	gb:BC006177.1 /DEF=Homo sapiens, Similar to metastasis associated 1, clone MGC:13258, mRNA, complete cds. /FEA=mRNA /PROD=Similar to metastasis associated 1 /DB_XREF=gi:13544097 /FL=gb:BC006177.1	BC006177		NP_004680
211796_s_at	0.00161	gb:AF043179.1 /DEF=Homo sapiens T cell receptor beta chain (TCRBV13S1-TCRBJ2S1) mRNA, complete cds. /FEA=CDS /GEN=TCRBV13S1-TCRBJ2S1 /PROD=T cell receptor beta chain /DB_XREF=gi:3002924 /UG=Hs.303157 T cell receptor beta locus /FL=gb:AF043179.1	AF043179		
211801_x_at	0.044231	gb:AF329637.1 /DEF=Homo sapiens mitofusin 1 mRNA, nuclear gene for mitochondrial protein, complete cds. /FEA=CDS /PROD=mitofusin 1 precursor /DB_XREF=gi:12744895 /UG=Hs.197877 hypothetical protein FLJ20693 /FL=gb:AF329637.1	AF329637		NP_284941
211814_s_at	0.012993	gb:AF112857.1 /DEF=Homo sapiens cyclin E2 splice variant 1 mRNA, complete cds. /FEA=CDS /PROD=cyclin E2 splice variant 1 /DB_XREF=gi:4092787 /UG=Hs.30464 cyclin E2 /FL=gb:AF112857.1	AF112857		NP_477097

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211816_x_at	0.004075	gb:D87858.1 /DEF=Homo sapiens mRNA for CD89_U04, complete cds. /FEA=CDS /PROD=CD89_U04 /DB_XREF=gi:1561545 /UG=Hs.193122 Fc fragment of IgA, receptor for /FL=gb:D87858.1 gb:D87860.1	D87858		NP_579814
211822_s_at	0.008036	gb:AF229061.1 /DEF=Homo sapiens NAC-gamma splice variant (NAC) mRNA, complete cds, alternatively spliced. /FEA=CDS /GEN=NAC /PROD=NAC-gamma splice variant /DB_XREF=gi:12656108 /UG=Hs.104305 death effector filament-forming Ced-4-like apoptosis protein /FL=gb:AF229061.1	AF229061		NP_127500
211825_s_at	0.04734	gb:AF327066.1 /DEF=Homo sapiens Ewings sarcoma EWS-Fli1 (type 1) oncogene mRNA, complete cds. /FEA=CDS /PROD=Ewings sarcoma EWS-Fli1 (type 1) oncogene /DB_XREF=gi:12963354 /UG=Hs.129953 Ewing sarcoma breakpoint region 1 /FL=gb:AF327066.1	AF327066		
211858_x_at	1.26E-04	gb:AF088184.1 /DEF=Homo sapiens guanine nucleotide-binding protein Gs alpha subunit isoform L2 mRNA, complete cds. /FEA=CDS /PROD=guanine nucleotide-binding protein Gs alphasubunit isoform L2 /DB_XREF=gi:3552031 /UG=Hs.273385 guanine nucleotide binding protein (G protein), alpha stimulating activity polypeptide 1 /FL=gb:AF088184.1	AF088184		NP_536351

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211883_x_at	0.041055	gb:M76742.1 /DEF=Homo sapiens alternatively spliced biliary glycoprotein (BGP _a) mRNA, complete cds. /FEA=CDS /GEN=BGP _a /PROD=biliary glycoprotein /DB_XREF=gi:179480 /UG=Hs.50964 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein) /FL=gb:M76742.1	M76742		NP_001703
211889_x_at	0.026891	gb:D12502.1 /DEF=Human mRNA for biliary glycoprotein, complete cds. /FEA=CDS /GEN=BGP /PROD=biliary glycoprotein /DB_XREF=gi:219494 /UG=Hs.50964 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein) /FL=gb:D12502.1	D12502		NP_001703
211902_x_at	0.004341	gb:L34703.1 /DEF=Homo sapiens T-cell receptor alpha chain (TCRA) mRNA (HLA-A1, 24; B7, 8; DR 1, 3), complete cds. /FEA=CDS /GEN=TCRA /PROD=T-cell receptor alpha /DB_XREF=gi:1100165 /UG=Hs.74647 Human T-cell receptor active alpha-chain mRNA from JM cell line, complete cds /FL=gb:L34703.1	L34703		
211909_x_at	0.046651	gb:L32662.1 /DEF=Human prostaglandin E2 receptor EP3 subtype isoform IV mRNA, complete cds. /FEA=CDS /PROD=prostaglandin E2 receptor /DB_XREF=gi:484163 /FL=gb:L32662.1	L32662		
211913_s_at	0.008391	gb:L08961.1 /DEF=Homo sapiens transmembrane tyrosine kinase mRNA, complete cds. /FEA=CDS /PROD=tyrosine kinase /DB_XREF=gi:897614 /FL=gb:L08961.1	L08961		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211921_x_at	0.033381	gb:AF348514.1 /DEF=Homo sapiens fetal thymus prothymosin alpha mRNA, complete cds. /FEA=CDS /PROD=prothymosin alpha /DB_XREF=gi:13560658 /FL=gb:AF348514.1	AF348514		NP_002814
211926_s_at	0.041055	myosin, heavy polypeptide 9, non-muscle	AI827941	Hs.146550	NP_002464
211927_x_at	0.02145	ESTs, Highly similar to S22655 translation elongation factor eEF-1 gamma chain - human [H.sapiens]	BE963164	Hs.381229	
211928_at	0.004341	Consensus includes gb:AB002323.1 /DEF=Human mRNA for KIAA0325 gene, partial cds. /FEA=mRNA /GEN=KIAA0325 /DB_XREF=gi:2224590 /UG=Hs.7720 dynein, cytoplasmic, heavy polypeptide 1	AB002323		NP_001367
211930_at	0.004341	Homo sapiens BX1 mRNA, partial cds	BE867771	Hs.249247	
211936_at	7.58E-04	Consensus includes gb:AF216292.1 /DEF=Homo sapiens endoplasmic reticulum lumenal Ca2+ binding protein grp78 mRNA, complete cds. /FEA=CDS /PROD=endoplasmic reticulum lumenal Ca2+ bindingprotein grp78 /DB_XREF=gi:7229461 /UG=Hs.75410 heat shock 70kD protein 5 (glucose-regulated protein, 78kD) /FL=gb:AF216292.1	AF216292		NP_005338
211937_at	0.003171	Consensus includes gb:NM_001417.1 /DEF=Homo sapiens eukaryotic translation initiation factor 4B (EIF4B), mRNA. /FEA=CDS /GEN=EIF4B /PROD=eukaryotic translation initiation factor 4B /DB_XREF=gi:4503532 /UG=Hs.93379 eukaryotic translation initiation factor 4B /FL=gb:NM_001417.1	NM_001417		NP_001408
211941_s_at	0.033381	prostatic binding protein	BF686267	Hs.80423	NP_002558

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
211944_at	0.003539	Consensus includes gb:BE729523 /FEA=EST /DB_XREF=gi:10143515 /DB_XREF=est:601564896F1 /CLONE=IMAGE:3926279 /UG=Hs.69559 KIAA1096 protein	AL096857		NP_055987
211946_s_at	0.026891	Consensus includes gb:AL096857.1 /DEF=Novel human mRNA from chromosome 1, which has similarities to BAT2 genes. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:5541862 /UG=Hs.69559 KIAA1096 protein	AL096857		NP_055987
211948_x_at	6.68E-04	Consensus includes gb:BG261071 /FEA=EST /DB_XREF=gi:12770887 /DB_XREF=est:602372693F1 /CLONE=IMAGE:4480631 /UG=Hs.69559 KIAA1096 protein	AL096857		NP_055987
211950_at	0.041055	Consensus includes gb:AB007931.1 /DEF=Homo sapiens mRNA for KIAA0462 protein, partial cds. /FEA=mRNA /GEN=KIAA0462 /PROD=KIAA0462 protein /DB_XREF=gi:3413885 /UG=Hs.297641 KIAA0462 protein	AB007931		NP_065816
211960_s_at	1.26E-04	Consensus includes gb:BG261416 /FEA=EST /DB_XREF=gi:12771232 /DB_XREF=est:602373192F1 /CLONE=IMAGE:4484422 /UG=Hs.237955 hypothetical protein PRO2706	AK000826		NP_004628
211961_s_at	1.26E-04	Consensus includes gb:AK000826.1 /DEF=Homo sapiens cDNA FLJ20819 fis, clone ADSE00511. /FEA=mRNA /DB_XREF=gi:7021139 /UG=Hs.237955 hypothetical protein PRO2706	AK000826		NP_004628
211963_s_at	5.03E-04	actin related protein 2/3 complex, subunit 5, 16kDa	AL516350	Hs.82425	NP_005708

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
211967_at	0.026891	pro-oncosis receptor inducing membrane injury gene	BG538627	Hs.393431	NP_443164
211974_x_at	0.026891	recombining binding protein suppressor of hairless (Drosophila)	AL513759	Hs.356806	
211976_at	5.03E-04	Consensus includes gb:AK026168.1 /DEF=Homo sapiens cDNA: FLJ22515 fis, clone HRC12122, highly similar to AF052101 Homo sapiens clone 23872 mRNA sequence. /FEA=mRNA /DB_XREF=gi:10438931 /UG=Hs.188882 Homo sapiens cDNA: FLJ21862 fis, clone HEP02321, highly similar to AF052101 Homo sapiens clone 23872 mRNA sequence	AK026168		
211984_at	0.00161	ESTs, Moderately similar to apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 2; apolipoprotein B mRNA editing enzyme, catalytic polypeptide 2 [Homo sapiens] [H.sapiens]	AI653730	Hs.374441	
211985_s_at	5.03E-04	ESTs, Moderately similar to apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 2; apolipoprotein B mRNA editing enzyme, catalytic polypeptide 2 [Homo sapiens] [H.sapiens]	AI653730	Hs.374441	
211993_at	0.027792	protein kinase, lysine deficient 1	AI742553	Hs.184592	NP_061852
211994_at	1.26E-04	protein kinase, lysine deficient 1	AI742553	Hs.184592	NP_061852
211997_x_at	0.033381	Consensus includes gb:NM_005324.1 /DEF=Homo sapiens H3 histone, family 3B (H3.3B) (H3F3B), mRNA. /FEA=CDS /GEN=H3F3B /PROD=H3 histone, family 3B (H3.3B) /DB_XREF=gi:4885384 /UG=Hs.180877 H3 histone, family 3B (H3.3B) /FL=gb:NM_005324.1	NM_005324		NP_005315
212006_at	0.033381	Consensus includes gb:AU149908 /FEA=EST /DB_XREF=gi:11011429 /DB_XREF=est:AU149908 /CLONE=NT2RP2000896 /UG=Hs.77495 UBX domain-containing 1	D87684		NP_055422

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212007_at	0.011626	Consensus includes gb:AI927512 /FEA=EST /DB_XREF=gi:5663476 /DB_XREF=est:wo90g11.x1 /CLONE=IMAGE:2462660 /UG=Hs.77495 UBX domain-containing 1	D87684		NP_055422
212010_s_at	0.001116	Consensus includes gb:AK025647.1 /DEF=Homo sapiens cDNA: FLJ21994 fis, clone HEP06577, highly similar to AF103803 Homo sapiens clone H41 unknown mRNA. /FEA=mRNA /DB_XREF=gi:10438231 /UG=Hs.283690 hypothetical protein /FL=gb:NM_017548.1	AK025647		NP_060018
212016_s_at	0.020793	polypyrimidine tract binding protein 1	AA679988	Hs.172550	NP_787041
212017_at	0.026891	ESTs, Weakly similar to S57447 HPBRII-7 protein - human [H.sapiens]	BF677404	Hs.409180	
212020_s_at	0.002537	antigen identified by monoclonal antibody Ki-67	BF001806	Hs.80976	NP_002408
212024_x_at	2.17E-04	see GenBank Accession Number U01184 for cDNA; similar to Drosophila melanogaster flil in GenBank Accession Number U01182 and Caenorhabditis elegans flil homolog in GenBank Accession Number U01183; Homo sapiens FLII gene, complete cds.	NM_002018		NP_002009
212025_s_at	0.001054	see GenBank Accession Number U01184 for cDNA; similar to Drosophila melanogaster flil in GenBank Accession Number U01182 and Caenorhabditis elegans flil homolog in GenBank Accession Number U01183; Homo sapiens FLII gene, complete cds.	NM_002018		NP_002009
212027_at	1.26E-04	Homo sapiens cDNA: FLJ22454 fis, clone HRC09703	BE466128	Hs.409075	
212034_s_at	0.004341	likely ortholog of mouse exocyst component protein 70 kDa homolog (S. cerevisiae) Exo70: exocyst component protein 70 kDa homolog (S. cerevisiae)	BE646386	Hs.325530	NP_056034
212039_x_at	0.033381	ribosomal protein L3	BG339228	Hs.119598	NP_000958

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212040_at	0.033381	Consensus includes gb:BG249599 /FEA=EST /DB_XREF=gi:12759426 /DB_XREF=est:602319692F1 /CLONE=IMAGE:4414895 /UG=Hs.110771 Homo sapiens cDNA: FLJ21904 fis, clone HEP03585	AK025557		
212043_at	3.26E-04	Consensus includes gb:W72053 /FEA=EST /DB_XREF=gi:1382323 /DB_XREF=est:zd66e04.s1 /CLONE=IMAGE:345630 /UG=Hs.110771 Homo sapiens cDNA: FLJ21904 fis, clone HEP03585	AK025557		
212044_s_at	0.001116	ribosomal protein L27a	BE737027	Hs.76064	NP_000981
212046_x_at	0.041055	Consensus includes gb:X60188.1 /DEF=Human ERK1 mRNA for protein serinethreonine kinase. /FEA=mRNA /GEN=ERK1 /PROD=protein serinethreonine kinase /DB_XREF=gi:31220 /UG=Hs.861 mitogen-activated protein kinase 3	X60188		
212051_at	2.05E-04	Consensus includes gb:AA676803 /FEA=EST /DB_XREF=gi:2657325 /DB_XREF=est:zj65b04.s1 /CLONE=IMAGE:455119 /UG=Hs.13996 Homo sapiens cDNA: FLJ23260 fis, clone COL05804, highly similar to HSU90911 Human clone 23652 mRNA sequence	AK026913		NP_573571
212052_s_at	0.026891	Consensus includes gb:AB014576.1 /DEF=Homo sapiens mRNA for KIAA0676 protein, partial cds. /FEA=mRNA /GEN=KIAA0676 /PROD=KIAA0676 protein /DB_XREF=gi:3327165 /UG=Hs.155829 KIAA0676 protein	AB014576		NP_055858
212055_at	1.26E-04	DKFZP586M1523 protein	BF689173	Hs.22981	NP_056291

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212057_at	1.26E-04	Consensus includes gb:AA206161 /FEA=EST /DB_XREF=gi:1801720 /DB_XREF=est:zq55f01.s1 /CLONE=IMAGE:645529 /UG=Hs.75909 KIAA0182 protein	D80004		
212060_at	0.041055	Consensus includes gb:AU152088 /FEA=EST /DB_XREF=gi:11013609 /DB_XREF=est:AU152088 /CLONE=NT2RP3000162 /UG=Hs.7976 KIAA0332 protein	AB002330		
212063_at	5.03E-04	ESTs, Weakly similar to TRHY_HUMAN Trichohyalin [H.sapiens]	BE903880	Hs.408878	NP_000601
212065_s_at	0.003395	Consensus includes gb:AW502434 /FEA=EST /DB_XREF=gi:7116911 /DB_XREF=est:UI-HF-BR0p-ajs-g-03-0-UI.r1 /CLONE=IMAGE:3075652 /UG=Hs.180948 KIAA0729 protein	AB018272		
212066_s_at	0.013223	Consensus includes gb:AB018272.1 /DEF=Homo sapiens mRNA for KIAA0729 protein, partial cds. /FEA=mRNA /GEN=KIAA0729 /PROD=KIAA0729 protein /DB_XREF=gi:3882178 /UG=Hs.180948 KIAA0729 protein	AB018272		
212071_s_at	7.58E-04	spectrin, beta, non-erythrocytic 1	BE968833	Hs.107164	NP_842565
212072_s_at	0.001116	casein kinase 2, alpha 1 polypeptide pseudogene	AI161318	Hs.375587	
212076_at	0.011626	Consensus includes gb:AI701430 /FEA=EST /DB_XREF=gi:4989330 /DB_XREF=est:we29h08.x1 /CLONE=IMAGE:2342559 /UG=Hs.199160 myeloidlymphoid or mixed-lineage leukemia (trithorax (Drosophila) homolog) /FL=gb:L04284.1 gb:NM_005933.1	NM_005933		NP_005924

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212080_at	0.022795	Consensus includes gb:AV714029 /FEA=EST /DB_XREF=gi:10795546 /DB_XREF=est:AV714029 /CLONE=DCBCDA03 /UG=Hs.199160 myeloidlymphoid or mixed-lineage leukemia (trithorax (Drosophila) homolog) /FL=gb:L04284.1 gb:NM_005933.1	NM_005933		NP_005924
212082_s_at	0.00228	myosin, light polypeptide 6, alkali, smooth muscle and non-muscle	BE734356	Hs.77385	NP_524149
212086_x_at	0.016686	Consensus includes gb:AK026584.1 /DEF=Homo sapiens cDNA: FLJ22931 fis, clone KAT07501, highly similar to HSLAMAR Human mRNA for nuclear envelope protein lamin A precursor. /FEA=mRNA /DB_XREF=gi:10439468 /UG=Hs.77886 lamin AC	M13452		NP_733822
212098_at	0.00228	ESTs	AL134724	Hs.408961	
212099_at	0.041055	ras homolog gene family, member B	AI263909	Hs.204354	
212101_at	0.013575	Consensus includes gb:AU154321 /FEA=EST /DB_XREF=gi:11015842 /DB_XREF=est:AU154321 /CLONE=NT2RP4000774 /UG=Hs.301553 karyopherin alpha 6 (importin alpha 7) /FL=gb:AF060543.1 gb:NM_012316.1	NM_012316		NP_036448
212102_s_at	0.013223	Consensus includes gb:AK002111.1 /DEF=Homo sapiens cDNA FLJ11249 fis, clone PLACE1008790, highly similar to Homo sapiens importin alpha 7 subunit mRNA. /FEA=mRNA /DB_XREF=gi:7023792 /UG=Hs.301553 karyopherin alpha 6 (importin alpha 7) /FL=gb:AF060543.1 gb:NM_012316.1	NM_012316		NP_036448

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212112_s_at	0.003171	Consensus includes gb:AI816243 /FEA=EST /DB_XREF=gi:5431789 /DB_XREF=est:au46a11.x1 /CLONE=IMAGE:2517788 /UG=Hs.106823 H.sapiens gene from PAC 426I6, similar to syntaxin 7	AL035306		NP_803173
212120_at	1.26E-04	ras-like protein TC10	BF348067	Hs.250697	NP_036381
212124_at	5.03E-04	Consensus includes gb:AF070622.1 /DEF=Homo sapiens clone 24800 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3283888 /UG=Hs.7252 KIAA1224 protein	AF070622		
212132_at	0.016934	Consensus includes gb:AL117499.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D1335 (from clone DKFZp434D1335); partial cds. /FEA=mRNA /GEN=DKFZp434D1335 /PROD=hypothetical protein /DB_XREF=gi:5912003 /UG=Hs.8258 DKFZP434D1335 protein	AL117499		NP_056393
212135_s_at	7.58E-04	ribosomal protein L35a	AW517686	Hs.287361	NP_001675
212136_at	0.005852	ribosomal protein L35a	AW517686	Hs.287361	NP_001675
212137_at	5.03E-04	hypothetical protein MGC19556	AV746402	Hs.356357	NP_291029
212150_at	0.033381	KIAA0143 protein	AA805651	Hs.84087	
212153_at	0.038535	Consensus includes gb:AB007930.1 /DEF=Homo sapiens mRNA for KIAA0461 peroteine, partial cds. /FEA=mRNA /GEN=KIAA0461 /PROD=KIAA0461 peroteine /DB_XREF=gi:3413883 /UG=Hs.107088 KIAA0461 protein	AB007930		NP_665739
212159_x_at	1.26E-04	Consensus includes gb:AI125280 /FEA=EST /DB_XREF=gi:3593794 /DB_XREF=est:qd92c05.x1 /CLONE=IMAGE:1736936 /UG=Hs.19121 adaptor-related protein complex 2, alpha 2 subunit	AB020706		NP_036437

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212162_at	0.026891	Consensus includes gb:AK022873.1 /DEF=Homo sapiens cDNA FLJ12811 fis, clone NT2RP2002475. /FEA=mRNA /DB_XREF=gi:10434518 /UG=Hs.9873 likely homolog of rat kinase D-interacting substance of 220 kDa; KIAA1250 protein	AB033076		
212163_at	0.004341	Consensus includes gb:AB033076.1 /DEF=Homo sapiens mRNA for KIAA1250 protein, partial cds. /FEA=mRNA /GEN=KIAA1250 /PROD=KIAA1250 protein /DB_XREF=gi:6330860 /UG=Hs.9873 likely homolog of rat kinase D-interacting substance of 220 kDa; KIAA1250 protein	AB033076		
212165_at	0.003171	Consensus includes gb:AF070537.1 /DEF=Homo sapiens clone 24606 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387894 /UG=Hs.17481 Homo sapiens clone 24606 mRNA sequence	AL133052		NP_612400
212166_at	2.76E-04	RAN binding protein 16	H38643	Hs.172685	NP_055839
212168_at	1.26E-04	Consensus includes gb:AL514547 /FEA=EST /DB_XREF=gi:12778041 /DB_XREF=est:AL514547 /CLONE=CL0BB004ZC07 (3 prime) /UG=Hs.180895 putative brain nuclearly-targeted protein	AB018308		NP_690051
212171_x_at	0.016426	vascular endothelial growth factor	H95344	Hs.73793	NP_003367
212175_s_at	0.007779	Consensus includes gb:AL513611 /FEA=EST /DB_XREF=gi:12777105 /DB_XREF=est:AL513611 /CLONE=XCL0BB001ZB07 (3 prime) /UG=Hs.171811 adenylate kinase 2	AK023758		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212177_at	0.003171	Consensus includes gb:AW081113 /FEA=EST /DB_XREF=gi:6036265 /DB_XREF=est:xc29c08.x1 /CLONE=IMAGE:2585678 /UG=Hs.18368 DKFZP564B0769 protein	AL080186		NP_116259
212180_at	1.26E-04	Consensus includes gb:AK000311.1 /DEF=Homo sapiens cDNA FLJ20304 fis, clone HEP06749. /FEA=mRNA /DB_XREF=gi:7020310 /UG=Hs.5613 Homo sapiens mRNA; cDNA DKFZp564E2222 (from clone DKFZp564E2222)	AK000311		NP_005198
212184_s_at	0.001116	Consensus includes gb:AL117407.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D2050 (from clone DKFZp434D2050); partial cds. /FEA=mRNA /GEN=DKFZp434D2050 /PROD=hypothetical protein /DB_XREF=gi:5911992 /UG=Hs.109727 TAK1-binding protein 2; KIAA0733 protein	AL117407		NP_663317
212191_x_at	0.026891	ribosomal protein L13	AW574664	Hs.180842	NP_150254
212192_at	0.004341	Consensus includes gb:AI718937 /FEA=EST /DB_XREF=gi:5036193 /DB_XREF=est:as50b04.x1 /CLONE=IMAGE:2320591 /UG=Hs.109438 Homo sapiens clone 24775 mRNA sequence	AF052169		NP_612453
212193_s_at	0.004644	likely ortholog of mouse Ia related protein	BE881529	Hs.6214	NP_056130
212198_s_at	5.03E-04	Consensus includes gb:AL515964 /FEA=EST /DB_XREF=gi:12779457 /DB_XREF=est:AL515964 /CLONE=CS0DA001YL22 (3 prime) /UG=Hs.79305 KIAA0255 gene product	AL049539		
212200_at	0.016934	Consensus includes gb:AK025933.1 /DEF=Homo sapiens cDNA: FLJ22280 fis, clone HRC03841. /FEA=mRNA /DB_XREF=gi:10438600 /UG=Hs.100729 KIAA0692 protein	AB014592		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212202_s_at	0.00228	Consensus includes gb:BG493972 /FEA=EST /DB_XREF=gi:13455486 /DB_XREF=est:602542252F1 /CLONE=IMAGE:4673316 /UG=Hs.16492 DKFZP564G2022 protein	AF132733		NP_056312
212204_at	2.05E-04	Consensus includes gb:AL049944.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G2022 (from clone DKFZp564G2022); partial cds. /FEA=mRNA /GEN=DKFZp564G2022 /PROD=hypothetical protein /DB_XREF=gi:4884189 /UG=Hs.16492 DKFZP564G2022 protein	AF132733		NP_056312
212208_at	7.58E-04	Consensus includes gb:AK023837.1 /DEF=Homo sapiens cDNA FLJ13775 fis, clone PLACE4000369, moderately similar to Homo sapiens thyroid hormone receptor-associated protein complex component TRAP240 mRNA. /FEA=mRNA /DB_XREF=gi:10435894 /UG=Hs.4084 KIAA1025 protein	AK023837		
212211_at	1.26E-04	gene trap ankyrin repeat	AI986295	Hs.12329	NP_115593
212219_at	0.003171	Consensus includes gb:D38521.1 /DEF=Human mRNA for KIAA0077 gene, partial cds. /FEA=mRNA /GEN=KIAA0077 /DB_XREF=gi:559329 /UG=Hs.112396 KIAA0077 protein	D38521		
212223_at	0.016934	Consensus includes gb:AI926544 /FEA=EST /DB_XREF=gi:5662508 /DB_XREF=est:wo46c12.x1 /CLONE=IMAGE:2458390 /UG=Hs.303154 popeye protein 3	AL117536		
212225_at	0.035763	putative translation initiation factor	AL516854	Hs.150580	NP_005792

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212228_s_at	0.012555	Consensus includes gb:AC004382 /DEF=Homo sapiens Chromosome 16 BAC clone CIT987SK-A-152E5 /FEA=mRNA_8 /DB_XREF=gi:3252819 /UG=Hs.4288 hypothetical protein DKFZp434K046	AC004382		
212229_s_at	6.68E-04	Consensus includes gb:AK001699.1 /DEF=Homo sapiens cDNA FLJ10837 fis, clone NT2RP4001260, highly similar to Homo sapiens mRNA for KIAA0875 protein. /FEA=mRNA /DB_XREF=gi:7023121 /UG=Hs.184227 F-box only protein 21 /FL=gb:AF174601.1	AK001699		NP_296373
212230_at	0.014233	ribosomal protein S20	AL576654	Hs.8102	NP_001014
212239_at	0.033381	Consensus includes gb:AI680192 /FEA=EST /DB_XREF=gi:4890374 /DB_XREF=est:tw66a10.x1 /CLONE=IMAGE:2264634 /UG=Hs.6241 phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)	M61906		NP_852556
212240_s_at	2.05E-04	Consensus includes gb:AI679268 /FEA=EST /DB_XREF=gi:4889450 /DB_XREF=est:tu62e04.x1 /CLONE=IMAGE:2255646 /UG=Hs.6241 phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)	M61906		NP_852556
212242_at	3.26E-04	tubulin, alpha 1 (testis specific)	AL565074	Hs.75318	NP_005991
212248_at	0.00228	Homo sapiens LYRIC mRNA, complete cds	AI972475	Hs.395896	
212249_at	0.008036	Consensus includes gb:AI934473 /FEA=EST /DB_XREF=gi:5673433 /DB_XREF=est:wp58d05.x1 /CLONE=IMAGE:2465961 /UG=Hs.6241 phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)	M61906		NP_852556
212250_at	1.26E-04	Homo sapiens LYRIC mRNA, complete cds	AI972475	Hs.395896	
212252_at	0.02145	calcium/calmodulin-dependent protein kinase kinase 2, beta	AA181179	Hs.108708	NP_757380

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212256_at	5.03E-04	putative UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase T9	BE906572	Hs.394578	NP_060010
212259_s_at	0.001618	hematopoietic PBX-interacting protein	BF344265	Hs.8068	NP_065385
212261_at	0.001116	trinucleotide repeat containing 15	AL045800	Hs.323317	
212262_at	1.26E-04	Consensus includes gb:AA149639 /FEA=EST /DB_XREF=gi:1720440 /DB_XREF=est:zl39c06.s1 /CLONE=IMAGE:504298 /UG=Hs.15020 homolog of mouse quaking QKI (KH domain RNA binding protein) /FL=gb:AF142419.1 gb:AF142422.1	AF142419		
212264_s_at	3.26E-04	Consensus includes gb:BE645850 /FEA=EST /DB_XREF=gi:9970161 /DB_XREF=est:7e77c03.x1 /CLONE=IMAGE:3288484 /UG=Hs.154978 KIAA0261 protein	D87450		
212265_at	5.03E-04	Consensus includes gb:AL031781 /DEF=Human DNA sequence from clone 51J12 on chromosome 6q26-27. Contains the 3 part of the alternatively spliced gene for the human orthologs of mouse QKI-7 and QKI-7B (KH Domain RNA Binding proteins) and zebrafish ZKQ-1 (Quaking protein homolog). Con... /FEA=mRNA_1 /DB_XREF=gi:4038570 /UG=Hs.15020 homolog of mouse quaking QKI (KH domain RNA binding protein) /FL=gb:AF142419.1 gb:AF142422.1	AF142419		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212268_at	1.26E-04	Consensus includes gb:NM_030666.1 /DEF=Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1 (SERPINB1), mRNA. /FEA=CDS /GEN=SERPINB1 /PROD=serine (or cysteine) proteinase inhibitor, cladeB (ovalbumin), member 1 /DB_XREF=gi:13489086 /UG=Hs.183583 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1 /FL=gb:NM_030666.1	NM_030666		NP_109591
212271_at	1.26E-04	mitogen-activated protein kinase 1	AA195999	Hs.324473	NP_620407
212273_x_at	3.26E-04	GNAS complex locus	AI591100	Hs.374523	NP_536351
212277_at	0.003171	Consensus includes gb:AB014547.1 /DEF=Homo sapiens mRNA for KIAA0647 protein, partial cds. /FEA=mRNA /GEN=KIAA0647 /PROD=KIAA0647 protein /DB_XREF=gi:3327107 /UG=Hs.141727 myotubularin related protein 4 /FL=gb:NM_004687.1 gb:AF264717.1	AB014547		NP_004678
212293_at	0.010205	Consensus includes gb:BG111260 /FEA=EST /DB_XREF=gi:12604766 /DB_XREF=est:602281701F1 /CLONE=IMAGE:4369057 /UG=Hs.12259 KIAA0630 protein	AB014530		NP_689909
212297_at	1.26E-04	hypothetical protein FLJ20986	BF218804	Hs.324507	NP_078800
212301_at	1.26E-04	Consensus includes gb:D87440.1 /DEF=Human mRNA for KIAA0252 gene, partial cds. /FEA=mRNA /GEN=KIAA0252 /DB_XREF=gi:1665770 /UG=Hs.83419 KIAA0252 protein	D87440		NP_055953

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212306_at	0.041055	Consensus includes gb:AI741784 /FEA=EST /DB_XREF=gi:5110072 /DB_XREF=est:wg22h09.x1 /CLONE=IMAGE:2365889 /UG=Hs.108614 KIAA0627 protein; Drosophila multiple asters (Mast)-like homolog 2	AB014527		
212312_at	1.32E-04	Consensus includes gb:AL117381 /DEF=Human DNA sequence from clone RP5-857M17 on chromosome 20 Contains ESTs, STSs, GSSs and CpG islands. Contains the 3 part of the ID1 gene for inhibitor of DNA binding 1 (dominant negative helix-loop-helix protein), a gene for a novel protein simil... /FEA=mRNA_1 /DB_XREF=gi:11493211 /UG=Hs.305890 BCL2-like 1	AL117381		
212314_at	0.041055	Consensus includes gb:AB018289.1 /DEF=Homo sapiens mRNA for KIAA0746 protein, partial cds. /FEA=mRNA /GEN=KIAA0746 /PROD=KIAA0746 protein /DB_XREF=gi:3882212 /UG=Hs.49500 KIAA0746 protein	AB018289		
212315_s_at	0.023856	Consensus includes gb:AA502912 /FEA=EST /DB_XREF=gi:2237879 /DB_XREF=est:ne42d10.s1 /CLONE=IMAGE:900019 /UG=Hs.56966 KIAA0906 protein	AB020713		NP_079199
212317_at	0.005852	Consensus includes gb:AK022910.1 /DEF=Homo sapiens cDNA FLJ12848 fis, clone NT2RP2003391, highly similar to Homo sapiens mRNA for nuclear transport receptor. /FEA=mRNA /DB_XREF=gi:10434575 /UG=Hs.69235 transportin-SR /FL=gb:NM_012470.1	NM_012470		NP_036602

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212320_at	0.02145	Consensus includes gb:BC001002.1 /DEF=Homo sapiens, clone IMAGE:3447696, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3447696) /DB_XREF=gi:12654358 /UG=Hs.179661 tubulin, beta polypeptide	BC001002		NP_821133
212322_at	0.00161	Consensus includes gb:BE999972 /FEA=EST /DB_XREF=gi:10700248 /DB_XREF=est:7h15b02.x1 /CLONE=IMAGE:3316011 /UG=Hs.186613 sphingosine-1-phosphate lyase 1 /FL=gb:AF144638.1	AF144638		NP_003892
212323_s_at	3.26E-04	Consensus includes gb:AA524553 /FEA=EST /DB_XREF=gi:2265481 /DB_XREF=est:ng45h12.s1 /CLONE=IMAGE:937799 /UG=Hs.194737 KIAA0453 protein	AB007922		
212330_at	1.26E-04	transcription factor Dp-1	R60866	Hs.79353	NP_009042
212332_at	0.020793	Consensus includes gb:BF110947 /FEA=EST /DB_XREF=gi:10940637 /DB_XREF=est:7n30f02.x1 /CLONE=IMAGE:3566330 /UG=Hs.79362 retinoblastoma-like 2 (p130) /FL=gb:NM_005611.1	NM_005611		NP_005602
212334_at	1.26E-04	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AW167793	Hs.409152	NP_002067
212335_at	1.26E-04	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AW167793	Hs.409152	NP_002067
212340_at	2.05E-04	general transcription factor IIA, 2, 12kDa	AA195936	Hs.76362	
212341_at	0.005852	general transcription factor IIA, 2, 12kDa	AA195936	Hs.76362	
212342_at	0.011725	general transcription factor IIA, 2, 12kDa	AA195936	Hs.76362	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212350_at	0.016934	Consensus includes gb:AB029031.1 /DEF=Homo sapiens mRNA for KIAA1108 protein, partial cds. /FEA=mRNA /GEN=KIAA1108 /PROD=KIAA1108 protein /DB_XREF=gi:5689552 /UG=Hs.278586 KIAA1108 protein	AB029031		NP_055988
212355_at	1.26E-04	HCDI protein	AI075450	Hs.7911	
212356_at	0.00945	HCDI protein	AI075450	Hs.7911	
212367_at	0.016934	Consensus includes gb:AI799061 /FEA=EST /DB_XREF=gi:5364533 /DB_XREF=est:we98a10.x1 /CLONE=IMAGE:2349114 /UG=Hs.6048 FEM-1 (C.elegans) homolog b /FL=gb:AF178632.1 gb:NM_015322.1 gb:AF204883.1	NM_015322		NP_056137
212368_at	0.013223	hypothetical protein FLJ13564	AA972711	Hs.173081	
212369_at	3.26E-04	zinc finger protein 384	AI264312	Hs.103315	NP_597733
212370_x_at	0.041055	Consensus includes gb:AL080183.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434O171 (from clone DKFZp434O171); partial cds. /FEA=mRNA /GEN=DKFZp434O171 /PROD=hypothetical protein /DB_XREF=gi:5262659 /UG=Hs.13273 KIAA0592 protein	AL080183		
212371_at	0.041055	Consensus includes gb:AL049397.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586C1019 (from clone DKFZp586C1019). /FEA=mRNA /DB_XREF=gi:4500188 /UG=Hs.12314 Homo sapiens mRNA; cDNA DKFZp586C1019 (from clone DKFZp586C1019)	AL049397		

Gene Identifier	p-value	Description	Gen Accession No.	Unigene Accession No.	Protein Accession No.
212373_at	0.001116	Consensus includes gb:AW139179 /FEA=EST /DB_XREF=gi:6143497 /DB_XREF=est:UI-H-BI1-aet-f-06-0-UI.s1 /CLONE=IMAGE:2720411 /UG=Hs.6048 FEM-1 (C.elegans) homolog b /FL=gb:AF178632.1 gb:Nm_015322.1 gb:AF204883.1	NM_015322		NP_056137
212377_s at	5.03E-04	Notch homolog 2 (Drosophila)	AU158495	Hs.131286	NP_077719
212382_at	0.029453	Consensus includes gb:BF433429 /FEA=EST /DB_XREF=gi:11445592 /DB_XREF=est:7q53a12.x1 /CLONE=IMAGE:3702046 /UG=Hs.289068 Homo sapiens cDNA FLJ11918 fis, clone HEMBB1000272	AK021980		
212383_at	0.005852	Consensus includes gb:AL096733.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434E071 (from clone DKFZp434E071). /FEA=mRNA /DB_XREF=gi:5419866 /UG=Hs.267871 ATPase, H+ transporting, lysosomal (vacuolar proton pump) non-catalytic accessory protein 1A (110116kD)	AL096733		NP_005168
212388_at	0.003862	Consensus includes gb:AB028980.1 /DEF=Homo sapiens mRNA for KIAA1057 protein, partial cds. /FEA=mRNA /GEN=KIAA1057 /PROD=KIAA1057 protein /DB_XREF=gi:5689450 /UG=Hs.7243 ubiquitin specific protease 24	AB028980		
212402_at	0.004341	KIAA0853 protein	BE895685	Hs.136102	NP_055885
212403_at	0.005852	Consensus includes gb:A1749193 /FEA=EST /DB_XREF=gi:5127457 /DB_XREF=est:at40e04.x1 /CLONE=IMAGE:2374494 /UG=Hs.17639 Homo sapiens ubiquitin protein ligase (UBE3B) mRNA, partial cds	AL096740		NP_569733

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212408_at	1.26E-04	Consensus includes gb:AK023204.1 /DEF=Homo sapiens cDNA FLJ13142 fis, clone NT2RP3003212, moderately similar to Rattus norvegicus lamina associated polypeptide 1C (LAP1C) mRNA. /FEA=mRNA /DB_XREF=gi:10435025 /UG=Hs.234265 DKFZP586G011 protein	AK023204		NP_056417
212412_at	0.003171	amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	AV715767	Hs.177486	
212421_at	0.00161	Consensus includes gb:AB023147.1 /DEF=Homo sapiens mRNA for KIAA0930 protein, partial cds. /FEA=mRNA /GEN=KIAA0930 /PROD=KIAA0930 protein /DB_XREF=gi:4589503 /UG=Hs.13255 KIAA0930 protein	AB023147		
212428_at	3.26E-04	Consensus includes gb:AW001101 /FEA=EST /DB_XREF=gi:5848017 /DB_XREF=est:wu24c05.x1 /CLONE=IMAGE:2520968 /UG=Hs.3852 KIAA0368 protein	AB002366		
212429_s_at	0.016686	general transcription factor IIIC, polypeptide 2, beta 110kDa	AW194657	Hs.75782	NP_001512
212433_x_at	0.005852	cadherin 1, type 1, E-cadherin (epithelial)	AA630314	Hs.356360	NP_002943
212434_at	0.016686	GrpE-like protein cochaperone	AL542571	Hs.151903	NP_079472
212436_at	7.44E-04	tripartite motif-containing 33	AI967961	Hs.287414	NP_148980
212437_at	0.033381	Consensus includes gb:AL109804 /DEF=Human DNA sequence from clone RP5-1009E24 on chromosome 20 Contains a novel gene encoding two isoforms similar to mouse sialoadhesin (a macrophage sialic acid binding receptor), a novel gene similar to KIAA0417, the CENPB gene (centromere protein ... /FEA=mRNA_8 /DB_XREF=gi:11121192 /UG=Hs.85004 centromere protein B (80kD)	AL109804		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212438_at	0.02145	putative nucleic acid binding protein RY-1	BG252325	Hs.54649	NP_006848
212440_at	0.013223	putative nucleic acid binding protein RY-1	BG252325	Hs.54649	NP_006848
212441_at	0.003171	Consensus includes gb:D86985.2 /DEF=Homo sapiens mRNA for KIAA0232 protein, partial cds. /FEA=mRNA /GEN=KIAA0232 /PROD=KIAA0232 protein /DB_XREF=gi:6634002 /UG=Hs.79276 KIAA0232 gene product	D86985		
212449_s_at	2.05E-04	lysophospholipase I	BG288007	Hs.409096	NP_006321
212451_at	0.018693	Consensus includes gb:N52532 /FEA=EST /DB_XREF=gi:1193698 /DB_XREF=est:yv35b11.s1 /CLONE=IMAGE:244701 /UG=Hs.118978 KIAA0256 gene product	D87445		
212452_x_at	8.68E-04	Consensus includes gb:AF113514.1 /DEF=Homo sapiens histone acetyltransferase MORF mRNA, complete cds. /FEA=CDS /PROD=histone acetyltransferase MORF /DB_XREF=gi:6002685 /UG=Hs.27590 histone acetyltransferase /FL=gb:AF113514.1	AF113514		NP_036462
212458_at	1.92E-04	ESTs, Weakly similar to SPY3_HUMAN Sprouty homolog 3 (Spry-3) [H.sapiens]	AW138902	Hs.392741	
212460_at	1.26E-04	hypothetical protein MGC24447	BE738425	Hs.353161	NP_612145
212461_at	1.26E-04	protease, serine, 15	BF793951	Hs.350265	NP_680479
212462_at	0.001618	Consensus includes gb:AU144267 /FEA=EST /DB_XREF=gi:11005788 /DB_XREF=est:AU144267 /CLONE=HEMBA1001396 /UG=Hs.27590 histone acetyltransferase /FL=gb:AF113514.1	AF113514		NP_036462
212463_at	0.004341	CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344)	BE379006	Hs.42346	
212465_at	0.010205	hypothetical protein FLJ23027	AA524500	Hs.321766	NP_115609

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212467_at	1.26E-04	Consensus includes gb:AB014578.1 /DEF=Homo sapiens mRNA for KIAA0678 protein, partial cds. /FEA=mRNA /GEN=KIAA0678 /PROD=KIAA0678 protein /DB_XREF=gi:3327169 /UG=Hs.12707 KIAA0678 protein	AB014578		
212468_at	2.05E-04	Consensus includes gb:AK023512.1 /DEF=Homo sapiens cDNA FLJ13450 fis, clone PLACE1003027, highly similar to Homo sapiens mRNA for KIAA0516 protein. /FEA=mRNA /DB_XREF=gi:10435467 /UG=Hs.129872 sperm associated antigen 9	AB011088		NP_758853
212472_at	0.001116	KIAA0750 gene product	BE965029	Hs.314434	
212473_s_at	0.013223	KIAA0750 gene product	BE965029	Hs.314434	
212474_at	0.004341	KIAA0241 protein	BE503381	Hs.150275	
212476_at	0.00228	Consensus includes gb:D26069.1 /DEF=Human mRNA for KIAA0041 gene, partial cds. /FEA=mRNA /GEN=KIAA0041 /DB_XREF=gi:436227 /UG=Hs.24340 centaurin beta2	D26069		NP_036419
212477_at	0.001303	Consensus includes gb:D26069.1 /DEF=Human mRNA for KIAA0041 gene, partial cds. /FEA=mRNA /GEN=KIAA0041 /DB_XREF=gi:436227 /UG=Hs.24340 centaurin beta2	D26069		NP_036419
212479_s_at	0.007779	Consensus includes gb:AK022815.1 /DEF=Homo sapiens cDNA FLJ12753 fis, clone NT2RP2001226. /FEA=mRNA /DB_XREF=gi:10434432 /UG=Hs.75277 hypothetical protein FLJ13910	AL050139		NP_073617
212481_s_at	5.03E-04	tropomyosin 4	AI214061	Hs.250641	NP_003281

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212483_at	0.026891	Consensus includes gb:AW339587 /FEA=EST /DB_XREF=gi:6836213 /DB_XREF=est:he14g08.x1 /CLONE=IMAGE:2919038 /UG=Hs.225767 IDN3 protein	AB019494		NP_597677
212491_s_at	0.027243	DnaJ (Hsp40) homolog, subfamily C, member 8	AA843895	Hs.74711	NP_055095
212492_s_at	0.003171	KIAA0876 protein	AW237172	Hs.301011	
212496_s_at	0.00228	KIAA0876 protein	AW237172	Hs.301011	
212503_s_at	0.026891	KIAA0934 protein	N31807	Hs.227716	
212506_at	2.05E-04	ubiquitin specific protease 2	AL135735	Hs.199160	NP_009097
212507_at	1.26E-04	Consensus includes gb:D87446.1 /DEF=Human mRNA for KIAA0257 gene, partial cds. /FEA=mRNA /GEN=KIAA0257 /DB_XREF=gi:1665780 /UG=Hs.75912 KIAA0257 protein	D87446		
212511_at	0.003171	ubiquitin specific protease 2	AL135735	Hs.199160	NP_009097
212513_s_at	0.026891	Consensus includes gb:AB029020.1 /DEF=Homo sapiens mRNA for KIAA1097 protein, partial cds. /FEA=mRNA /GEN=KIAA1097 /PROD=KIAA1097 protein /DB_XREF=gi:5689530 /UG=Hs.173694 KIAA1097 protein	AB029020		NP_055832
212517_at	0.026891	Consensus includes gb:AL132773 /DEF=Human DNA sequence from clone RP4-741H3 on chromosome 20 Contains parts of isoform 1 and isoform 2 (KIAA0548) of the ATRN (attractin (with dipeptidylpeptidase IV activity)) gene, ESTs, STSs and GSSs /FEA=mRNA_1 /DB_XREF=gi:7159777 /UG=Hs.194019 attractin	AL132773		
212519_at	0.00161	ubiquitin-conjugating enzyme E2E 1 (UBC4/5 homolog, yeast)	AL518159	Hs.7766	NP_003332

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212527_at	3.26E-04	Consensus includes gb:BF057059 /FEA=EST /DB_XREF=gi:10810955 /DB_XREF=est:7k15c07.x1 /CLONE=IMAGE:3444468 /UG=Hs.12421 hypothetical protein	AL023553		
212531_at	3.55E-04	Consensus includes gb:NM_005564.1 /DEF=Homo sapiens lipocalin 2 (oncogene 24p3) (LCN2), mRNA. /FEA=CDS /GEN=LCN2 /PROD=lipocalin 2 (oncogene 24p3) /DB_XREF=gi:5031852 /UG=Hs.204238 lipocalin 2 (oncogene 24p3) /FL=gb:NM_005564.1	NM_005564		NP_005555
212535_at	3.26E-04	MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A)	AA142929	Hs.277806	NP_005578
212536_at	1.26E-04	Consensus includes gb:AB023173.1 /DEF=Homo sapiens mRNA for KIAA0956 protein, partial cds. /FEA=mRNA /GEN=KIAA0956 /PROD=KIAA0956 protein /DB_XREF=gi:4589555 /UG=Hs.75478 ATPase, Class VI, type 11B	AB023173		
212542_s_at	0.041055	pleckstrin homology domain interacting protein	BF224151	Hs.367694	NP_060404
212543_at	1.26E-04	Consensus includes gb:U83115.1 /DEF=Human non-lens beta gamma-crystallin like protein (AIM1) mRNA, partial cds. /FEA=mRNA /GEN=AIM1 /PROD=non-lens beta gamma-crystallin like protein /DB_XREF=gi:2072424 /UG=Hs.161002 absent in melanoma 1	U83115		
212549_at	7.58E-04	Consensus includes gb:BE645861 /FEA=EST /DB_XREF=gi:9970172 /DB_XREF=est:7e77d05.x1 /CLONE=IMAGE:3288489 /UG=Hs.24064 Homo sapiens mRNA; cDNA DKFZp586N1323 (from clone DKFZp586N1323)	AL080218		
212552_at	0.001116	hippocalcin-like 1	BE617588	Hs.3618	NP_602293

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212553_at	0.033381	Consensus includes gb:AK026120.1 /DEF=Homo sapiens cDNA: FLJ22467 fis, clone HRC10345. /FEA=mRNA /DB_XREF=gi:10438867 /UG=Hs.29956 KIAA0460 protein	AK026120		
212560_at	0.016934	cytochrome c oxidase subunit VIb	AV728268	Hs.174031	
212561_at	0.003171	RAB6 interacting protein 1	AA349595	Hs.26797	
212570_at	0.009652	KIAA0830 protein	AL573201	Hs.167115	
212572_at	3.26E-04	Consensus includes gb:AW779556 /FEA=EST /DB_XREF=gi:7794159 /DB_XREF=est:hn81f05.x1 /CLONE=IMAGE:3034305 /UG=Hs.184523 KIAA0965 protein	AB023182		NP_055815
212583_at	0.011788	Consensus includes gb:AB011132.2 /DEF=Homo sapiens mRNA for KIAA0560 protein, partial cds. /FEA=mRNA /GEN=KIAA0560 /PROD=KIAA0560 protein /DB_XREF=gi:6635202 /UG=Hs.129952 KIAA0560 gene product	AB011132		
212584_at	0.010205	Consensus includes gb:BG260519 /FEA=EST /DB_XREF=gi:12770335 /DB_XREF=est:602372065F1 /CLONE=IMAGE:4480024 /UG=Hs.129952 KIAA0560 gene product	AB011132		
212586_at	0.02145	calpastatin	BG111635	Hs.9347	NP_775085
212590_at	7.58E-04	related RAS viral (r-ras) oncogene homolog 2	BG168858	Hs.206097	NP_036382
212594_at	0.033381	programmed cell death 4 (neoplastic transformation inhibitor)	N92498	Hs.326248	NP_663314
212597_s_at	1.26E-04	Consensus includes gb:AL079310.1 /DEF=Novel human gene mapping to chromosome 22. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:5102629 /UG=Hs.92260 high-mobility group protein 2-like 1	AL079310		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212599_at	0.00161	Consensus includes gb:AK025298.1 /DEF=Homo sapiens cDNA: FLJ21645 fis, clone COL08436. /FEA=mRNA /DB_XREF=gi:10437785 /UG=Hs.32168 KIAA0442 protein	AK025298		NP_056385
212601_at	0.004341	Consensus includes gb:AB007859.2 /DEF=Homo sapiens mRNA for KIAA0399 protein, partial cds. /FEA=mRNA /GEN=KIAA0399 /PROD=KIAA0399 protein /DB_XREF=gi:6634028 /UG=Hs.100955 KIAA0399 protein	AB007859		NP_055928
212602_at	2.05E-04	ALFY	AI806395	Hs.198135	NP_848700
212604_at	1.26E-04	Consensus includes gb:AI937794 /FEA=EST /DB_XREF=gi:5676664 /DB_XREF=est:wp82g10.x1 /CLONE=IMAGE:2468322 /UG=Hs.154655 imogen 38 /FL=gb:NM_005830.1	NM_005830		NP_005821
212606_at	1.26E-04	ALFY	AI806395	Hs.198135	NP_848700
212607_at	0.033381	Consensus includes gb:N32526 /FEA=EST /DB_XREF=gi:1152925 /DB_XREF=est:yy11f04.s1 /CLONE=IMAGE:270943 /UG=Hs.300642 serologically defined colon cancer antigen 8	U79271		NP_006633
212608_s_at	0.004341	ESTs, Moderately similar to Z254_HUMAN Zinc finger protein 254 (Bone marrow zinc finger 5) (BMZF-5) (Hematopoietic cell derived zinc finger protein 1) (HD-ZNF1) [H.sapiens]	W85912	Hs.409038	
212610_at	3.26E-04	Consensus includes gb:U79291.1 /DEF=Human clone 23721 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710271 /UG=Hs.83572 Human clone 23721 mRNA sequence	U79291		NP_116293

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212612_at	1.26E-04	Consensus includes gb:D31888.1 /DEF=Human mRNA for KIAA0071 gene, partial cds. /FEA=mRNA /GEN=KIAA0071 /DB_XREF=gi:506340 /UG=Hs.78398 KIAA0071 protein /FL=gb:AF155595.1 gb:NM_015156.1	D31888		NP_055971
212613_at	0.002418	butyrophilin, subfamily 3, member A2	AI991252	Hs.167741	NP_008978
212623_at	0.001116	Consensus includes gb:AU153138 /FEA=EST /DB_XREF=gi:11014659 /DB_XREF=est:AU153138 /CLONE=NT2RP3002507 /UG=Hs.174905 KIAA0033 protein	D26067		
212629_s_at	0.001054	Consensus includes gb:AI633689 /FEA=EST /DB_XREF=gi:4685019 /DB_XREF=est:th71d03.x1 /CLONE=IMAGE:2124101 /UG=Hs.69171 protein kinase C-like 2	AK023692		
212631_at	0.010205	Consensus includes gb:AI566082 /FEA=EST /DB_XREF=gi:4524534 /DB_XREF=est:tn53b12.x1 /CLONE=IMAGE:2172095 /UG=Hs.8906 Homo sapiens clone 24889 mRNA sequence	AF131808		NP_003560
212632_at	0.016934	Consensus includes gb:N32035 /FEA=EST /DB_XREF=gi:1152434 /DB_XREF=est:yw96e08.s1 /CLONE=IMAGE:260102 /UG=Hs.8906 Homo sapiens clone 24889 mRNA sequence	AF131808		NP_003560
212634_at	0.001618	KIAA0776 protein	AW298092	Hs.5460	NP_056138
212635_at	0.010205	ESTs	AW161626	Hs.409034	
212638_s_at	2.05E-04	WW domain-containing protein 1	BF131791	Hs.409154	NP_008944
212639_x_at	5.03E-04	tubulin, alpha, ubiquitous	AL581768	Hs.334842	NP_006073

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212642_s_at	0.041055	Consensus includes gb:AL023584 /DEF=Human DNA sequence from clone 67K17 on chromosome 6q24.1-24.3. Contains the HIVP2 (Schnurri-2) gene for HIV type 1 Enhancer-binding Protein 2, and a possible pseudogene in an intron of this gene. Contains STSs and GSSs and an AAAT repeat polymorp... /FEA=mRNA /DB_XREF=gi:3790154 /UG=Hs.75063 human immunodeficiency virus type 1 enhancer-binding protein 2 /FL=gb:NM_006734.1	AL023584		
212644_s_at	7.58E-04	likely ortholog of mouse MAPK-interacting and spindle-stabilizing protein	AI671747	Hs.388281	NP_653179
212649_at	0.004341	Consensus includes gb:AL079292.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 48814. /FEA=mRNA /PROD=hypothetical protein, similar to (AC007017)putative RNA helicase A Arabidopsis thaliana /DB_XREF=gi:5102732 /UG=Hs.95665 hypothetical protein	AL079292		NP_061903
212657_s_at	0.00161	interleukin 1 receptor antagonist	AW083357	Hs.81134	NP_776215
212658_at	0.00945	lipoma HMGIC fusion partner-like 2	N66633	Hs.79299	
212660_at	1.26E-04	KIAA0239 protein	AI735639	Hs.9729	NP_056103
212663_at	0.001116	Consensus includes gb:AB014574.1 /DEF=Homo sapiens mRNA for KIAA0674 protein, partial cds. /FEA=mRNA /GEN=KIAA0674 /PROD=KIAA0674 protein /DB_XREF=gi:3327161 /UG=Hs.14799 KIAA0674 protein	AB014574		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212666_at	7.58E-04	Consensus includes gb:AB046845.1 /DEF=Homo sapiens mRNA for KIAA1625 protein, partial cds. /FEA=mRNA /GEN=KIAA1625 /PROD=KIAA1625 protein /DB_XREF=gi:10047326 /UG=Hs.119120 E3 ubiquitin ligase SMURF1	AB046845		NP_851994
212669_at	0.003171	calcium/calmodulin-dependent protein kinase (CaM kinase) II gamma	AI093569	Hs.153406	NP_751913
212671_s_at	0.003171	major histocompatibility complex, class II, DQ alpha 1	BG397856	Hs.198253	NP_002113
212674_s_at	5.03E-04	Consensus includes gb:AK002076.1 /DEF=Homo sapiens cDNA FLJ11214 fis, clone PLACE1007990. /FEA=mRNA /DB_XREF=gi:7023738 /UG=Hs.281616 Homo sapiens cDNA FLJ11214 fis, clone PLACE1007990	AK002076		NP_619520
212677_s_at	0.004341	Consensus includes gb:BG530481 /FEA=EST /DB_XREF=gi:13522018 /DB_XREF=est:602559046F1 /CLONE=IMAGE:4697218 /UG=Hs.79507 KIAA0582 protein	AB011154		NP_055962
212678_at	0.011377	neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)	AW293356	Hs.93207	
212683_at	3.56E-04	KIAA0446 gene product	AL526243	Hs.158286	
212685_s_at	3.26E-04	Consensus includes gb:AI608789 /FEA=EST /DB_XREF=gi:4617956 /DB_XREF=est:tw94g12.x1 /CLONE=IMAGE:2267398 /UG=Hs.52515 transducin (beta)-like 2	AK026529		NP_116783

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212687_at	0.013223	Consensus includes gb:AL110164.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586I0324 (from clone DKFZp586I0324). /FEA=mRNA /DB_XREF=gi:5817069 /UG=Hs.193700 Homo sapiens mRNA; cDNA DKFZp586I0324 (from clone DKFZp586I0324)	AL110164		
212688_at	1.26E-04	Consensus includes gb:BC003393.1 /DEF=Homo sapiens, clone IMAGE:3447173, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3447173) /DB_XREF=gi:13097275 /UG=Hs.239818 phosphoinositide-3-kinase, catalytic, beta polypeptide /FL=gb:NM_006219.1	BC003393		NP_006210
212689_s_at	0.007779	Human putative zinc finger protein mRNA	AA524505	Hs.383008	NP_060903
212697_at	0.010205	Lutheran blood group (Auberger b antigen included)	AL515874	Hs.155048	NP_835227
212698_s_at	0.02008	hypothetical protein FLJ11619	BF966021	Hs.79844	NP_848699
212702_s_at	1.26E-04	Consensus includes gb:N45111 /FEA=EST /DB_XREF=gi:1186277 /DB_XREF=est:yz12f12.s1 /CLONE=IMAGE:282863 /UG=Hs.330988 Homo sapiens, Similar to Bicaudal D (Drosophila) homolog 1, clone IMAGE:3622452, mRNA, partial cds	BC004296		NP_056065
212710_at	0.025245	DKFZP434F195 protein	BE222901	Hs.184793	
212717_at	5.03E-04	Consensus includes gb:AB002354.2 /DEF=Homo sapiens mRNA for KIAA0356 protein, partial cds. /FEA=mRNA /GEN=KIAA0356 /PROD=KIAA0356 protein /DB_XREF=gi:6634022 /UG=Hs.32312 KIAA0356 gene product	AJ002220		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212722_s_at	0.026891	Consensus includes gb:AK021780.1 /DEF=Homo sapiens cDNA FLJ11718 fis, clone HEMBA1005252, highly similar to Homo sapiens mRNA for KIAA0585 protein. /FEA=mRNA /DB_XREF=gi:10433034 /UG=Hs.72660 phosphatidylserine receptor	AK021780		NP_055982
212723_at	1.26E-04	Consensus includes gb:AK021780.1 /DEF=Homo sapiens cDNA FLJ11718 fis, clone HEMBA1005252, highly similar to Homo sapiens mRNA for KIAA0585 protein. /FEA=mRNA /DB_XREF=gi:10433034 /UG=Hs.72660 phosphatidylserine receptor	AK021780		NP_055982
212725_s_at	0.026891	ESTs	N37081	Hs.409222	
212726_at	0.00161	Consensus includes gb:AB014562.1 /DEF=Homo sapiens mRNA for KIAA0662 protein, partial cds. /FEA=mRNA /GEN=KIAA0662 /PROD=KIAA0662 protein /DB_XREF=gi:3327137 /UG=Hs.93868 KIAA0662 gene product	AB014562		NP_078793
212737_at	0.00161	GM2 ganglioside activator protein	AL513583	Hs.289082	NP_000396
212740_at	0.001116	phosphoinositide-3-kinase, regulatory subunit 4, p150	BF740111	Hs.83050	NP_055417
212742_at	1.26E-04	zinc finger protein 364	AL530462	Hs.16537	
212743_at	0.007781	Consensus includes gb:AV694093 /FEA=EST /DB_XREF=gi:10295956 /DB_XREF=est:AV694093 /CLONE=GKCDSD10 /UG=Hs.48297 DKFZP586C1620 protein	AL050144		NP_056251
212744_at	0.002671	Bardet-Biedl syndrome 4	AI813772	Hs.387416	NP_149017
212747_at	1.28E-04	KIAA0229 protein	AI990523	Hs.221332	
212749_s_at	1.26E-04	Consensus includes gb:AI096477 /FEA=EST /DB_XREF=gi:3445971 /DB_XREF=est:qa03c06.x1 /CLONE=IMAGE:1685674 /UG=Hs.48297 DKFZP586C1620 protein	AL050144		NP_056251

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212752_at	7.58E-04	cytoplasmic linker associated protein 1	AA176798	Hs.392661	
212757_s_at	0.033381	calcium/calmodulin-dependent protein kinase (CaM kinase) II gamma	BF111268	Hs.153406	NP_751913
212761_at	0.016074	transcription factor 7-like 2 (T-cell specific, HMG-box)	AI703074	Hs.348412	NP_110383
212768_s_at	7.31E-04	isoform 1 match: proteins: Sw:Q07081 Tr:O95362 Tr:Q9Z2Y4 Tr:O95897 Tr:O70624 Sw:Q99972 Sw:Q99784 Sw:Q62609 Tr:Q9TV76 Tr:Q9I9K5 Sw:P01813 Tr:Q9IAK4 Tr:O35429; Human DNA sequence from clone RP11-209J19 on chromosome 13 Contains ESTs, STSs and GSSs. Contains the gene for the GW112 protein with two isoforms (GW112 and KIAA4294), complete sequence.	AL390736		
212769_at	0.026891	transducin-like enhancer of split 3 (E(sp1) homolog, Drosophila)	AI567426	Hs.156183	NP_005069
212779_at	0.004341	Consensus includes gb:AB029032.1 /DEF=Homo sapiens mRNA for KIAA1109 protein, partial cds. /FEA=mRNA /GEN=KIAA1109 /PROD=KIAA1109 protein /DB_XREF=gi:5689554 /UG=Hs.6606 KIAA1109 protein	AB029032		NP_115578
212780_at	0.007779	Consensus includes gb:AA700167 /FEA=EST /DB_XREF=gi:2703130 /DB_XREF=est:zj36h12.s1 /CLONE=IMAGE:452423 /UG=Hs.326392 son of sevenless (Drosophila) homolog 1 /FL=gb:L13857.1	L13857		NP_005624
212783_at	1.26E-04	Consensus includes gb:AI538172 /FEA=EST /DB_XREF=gi:4452307 /DB_XREF=est:ti75f08.x1 /CLONE=IMAGE:2137863 /UG=Hs.91065 hypothetical protein DKFZp761B2423	AK026954		NP_008841
212786_at	0.016934	KIAA0350 protein	AA731693	Hs.23263	
212787_at	3.26E-04	ZAP3 protein	AI952986	Hs.159471	
212790_x_at	0.010205	ribosomal protein L13a	BF942308	Hs.119122	NP_775866

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212794_s_at	0.013223	Consensus includes gb:AK001728.1 /DEF=Homo sapiens cDNA FLJ10866 fis, clone NT2RP4001614. /FEA=mRNA /DB_XREF=gi:7023169 /UG=Hs.12144 KIAA1033 protein	AL137753		
212796_s_at	0.041055	KIAA1055 protein	BF195608	Hs.126084	NP_055894
212799_at	3.26E-04	Consensus includes gb:BE217875 /FEA=EST /DB_XREF=gi:8905193 /DB_XREF=est:hv31a03.x1 /CLONE=IMAGE:3174988 /UG=Hs.12311 Homo sapiens clone 23570 mRNA sequence	AF038202		NP_005810
212802_s_at	7.58E-04	Consensus includes gb:AK023841.1 /DEF=Homo sapiens cDNA FLJ13779 fis, clone PLACE4000445, highly similar to Homo sapiens mRNA; cDNA DKFZp434C212 (from clone DKFZp434C212). /FEA=mRNA /DB_XREF=gi:10435900 /UG=Hs.172069 DKFZP434C212 protein	AK023841		
212804_s_at	0.026891	Consensus includes gb:AI797397 /FEA=EST /DB_XREF=gi:5362869 /DB_XREF=est:we87f12.x1 /CLONE=IMAGE:2348111 /UG=Hs.172069 DKFZP434C212 protein	AK023841		
212807_s_at	3.26E-04	sortilin 1	BE742268	Hs.351872	NP_002950
212808_at	0.041055	hypothetical protein FLJ14639	AA152202	Hs.334788	NP_116204
212811_x_at	5.03E-04	solute carrier family 1 (glutamate/neutral amino acid transporter), member 4	BG032165	Hs.323878	NP_003029
212814_at	0.033381	Consensus includes gb:AB020635.1 /DEF=Homo sapiens mRNA for KIAA0828 protein, partial cds. /FEA=mRNA /GEN=KIAA0828 /PROD=KIAA0828 protein /DB_XREF=gi:4240144 /UG=Hs.4984 KIAA0828 protein	AB020635		NP_056143
212823_s_at	0.007052	KIAA0599 protein	AU147160	Hs.198037	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212827_at	1.26E-04	Consensus includes gb:X17115.1 /DEF=Human mRNA for IgM heavy chain complete sequence. /FEA=mRNA /DB_XREF=gi:33450 /UG=Hs.302063 immunoglobulin heavy constant mu	X17115		
212828_at	0.009199	Consensus includes gb:AA191573 /FEA=EST /DB_XREF=gi:1780272 /DB_XREF=est:zp81b09.s1 /CLONE=IMAGE:626585 /UG=Hs.61289 synaptojanin 2	AL157424		NP_003889
212829_at	0.010205	ESTs	BE878277	Hs.57079	
212830_at	2.05E-04	EGF-like-domain, multiple 5	BF110421	Hs.5599	
212833_at	0.007779	Consensus includes gb:M74089.1 /DEF=Human TB1 gene mRNA, 3 end. /FEA=mRNA /GEN=TB1 /DB_XREF=gi:182400 /UG=Hs.75639 Human TB1 gene mRNA, 3 end	M74089		NP_620128
212836_at	0.016934	Consensus includes gb:D26018.1 /DEF=Human mRNA for KIAA0039 gene, partial cds. /FEA=mRNA /GEN=KIAA0039 /DB_XREF=gi:436221 /UG=Hs.82502 polymerase (DNA directed), delta 3	D26018		
212838_at	0.004452	Consensus includes gb:AB023227.1 /DEF=Homo sapiens mRNA for KIAA1010 protein, partial cds. /FEA=mRNA /GEN=KIAA1010 /PROD=KIAA1010 protein /DB_XREF=gi:4589669 /UG=Hs.23860 KIAA1010 protein	AB023227		
212840_at	0.016934	Consensus includes gb:BG339560 /FEA=EST /DB_XREF=gi:13145998 /DB_XREF=est:602437413F1 /CLONE=IMAGE:4555466 /UG=Hs.127287 KIAA0794 protein	AB018337		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212854_x_at	0.003171	Consensus includes gb:AB051480.1 /DEF=Homo sapiens mRNA for KIAA1693 protein, partial cds. /FEA=mRNA /GEN=KIAA1693 /PROD=KIAA1693 protein /DB_XREF=gi:12697930 /UG=Hs.323463 KIAA1693 protein; hypothetical protein FLJ20719	AB051480		NP_060410
212856_at	1.26E-04	Consensus includes gb:AB018310.1 /DEF=Homo sapiens mRNA for KIAA0767 protein, partial cds. /FEA=mRNA /GEN=KIAA0767 /PROD=KIAA0767 protein /DB_XREF=gi:3882254 /UG=Hs.95180 KIAA0767 protein	AB018310		
212861_at	0.003171	hypothetical protein MGC11308	BF690150	Hs.19210	NP_116278
212862_at	0.016934	CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2	AL568982	Hs.24812	NP_003809
212864_at	0.00161	CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2	AL568982	Hs.24812	NP_003809
212867_at	0.001116	nuclear receptor coactivator 2	AI040324	Hs.29131	NP_006531
212869_x_at	0.026891	tumor protein, translationally-controlled 1	AI721229	Hs.279860	NP_003286
212870_at	1.26E-04	son of sevenless homolog 2 (Drosophila)	AI628605	Hs.348496	
212871_at	0.020576	Consensus includes gb:NM_003668.1 /DEF=Homo sapiens mitogen-activated protein kinase-activated protein kinase 5 (MAPKAPK5), mRNA. /FEA=CDS /GEN=MAPKAPK5 /PROD=mitogen-activated protein kinase-activated protein kinase 5 /DB_XREF=gi:4506034 /UG=Hs.30327 mitogen-activated protein kinase-activated protein kinase 5 /FL=gb:NM_003668.1	NM_003668		NP_620777

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212872_s_at	0.005852	Consensus includes gb:AK023092.1 /DEF=Homo sapiens cDNA FLJ13030 fis, clone NT2RP3001111, highly similar to Homo sapiens TRF-proximal protein mRNA. /FEA=mRNA /DB_XREF=gi:10434851 /UG=Hs.93796 DKFZP586D2223 protein	AK023092		NP_061031
212880_at	0.041055	Consensus includes gb:AB011113.1 /DEF=Homo sapiens mRNA for KIAA0541 protein, partial cds. /FEA=mRNA /GEN=KIAA0541 /PROD=KIAA0541 protein /DB_XREF=gi:3043605 /UG=Hs.10881 WD repeat domain 7	AB011113		NP_443066
212882_at	3.26E-04	Consensus includes gb:AB018338.1 /DEF=Homo sapiens mRNA for KIAA0795 protein, partial cds. /FEA=mRNA /GEN=KIAA0795 /PROD=KIAA0795 protein /DB_XREF=gi:3882310 /UG=Hs.22926 KIAA0795 protein	AB018338		NP_079286
212887_at	1.26E-04	Sec23 homolog A (S. cerevisiae)	AI753659	Hs.272927	NP_006355
212888_at	0.007779	ESTs, Highly similar to A32915 nucleophosmin - human [H.sapiens]	BG109746	Hs.409325	
212890_at	0.003171	hypothetical protein MGC15523	BF663461	Hs.381200	NP_612637
212895_s_at	7.58E-04	active BCR-related gene	AL527773	Hs.408408	NP_068781
212899_at	7.58E-04	Consensus includes gb:AB028951.1 /DEF=Homo sapiens mRNA for KIAA1028 protein, partial cds. /FEA=mRNA /GEN=KIAA1028 /PROD=KIAA1028 protein /DB_XREF=gi:5689392 /UG=Hs.129836 KIAA1028 protein	AL122055		NP_055891

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212908_at	0.037567	Consensus includes gb:AK022530.1 /DEF=Homo sapiens cDNA FLJ12468 fis, clone NT2RM1000857, highly similar to Homo sapiens mRNA for KIAA0962 protein. /FEA=mRNA /DB_XREF=gi:10433971 /UG=Hs.9059 KIAA0962 protein	AB023179		
212911_at	0.041055	Consensus includes gb:AB023179.1 /DEF=Homo sapiens mRNA for KIAA0962 protein, partial cds. /FEA=mRNA /GEN=KIAA0962 /PROD=KIAA0962 protein /DB_XREF=gi:4589567 /UG=Hs.9059 KIAA0962 protein	AB023179		
212916_at	5.03E-04	KIAA1111 protein	AW249934	Hs.285641	
212917_x_at	0.003171	RecQ protein-like (DNA helicase Q1-like)	BF219234	Hs.235069	NP_079130
212920_at	0.010205	ESTs	AV682285	Hs.227630	
212926_at	0.008036	Consensus includes gb:AW183677 /FEA=EST /DB_XREF=gi:6452191 /DB_XREF=est:xj86e07.x1 /CLONE=IMAGE:2664132 /UG=Hs.103283 KIAA0594 protein	AB011166		NP_055925
212929_s_at	0.016686	hypothetical protein FLJ10824	W68158	Hs.13273	
212931_at	0.00228	Consensus includes gb:AB006630.1 /DEF=Homo sapiens mRNA for KIAA0292 gene, partial cds. /FEA=mRNA /GEN=KIAA0292 /DB_XREF=gi:2564331 /UG=Hs.201668 transcription factor 20 (AR1)	AB006630		
212932_at	0.013088	Consensus includes gb:AK022494.1 /DEF=Homo sapiens cDNA FLJ12432 fis, clone NT2RM1000018, highly similar to Human mRNA for KIAA0066 gene. /FEA=mRNA /DB_XREF=gi:10433912 /UG=Hs.227881 RAB3 GTPase-ACTIVATING PROTEIN	AK022494		
212934_at	1.26E-04	ESTs, Weakly similar to p47 [Homo sapiens] [H.sapiens]	AI245523	Hs.131357	
212936_at	0.013223	hypothetical protein DKFZp564D172	AI927701	Hs.25307	NP_114431

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212947_at	0.007779	Consensus includes gb:AL031685 /DEF=Human DNA sequence from clone RP5-963K23 on chromosome 20q13.11-13.2 Contains a KRT18 (Keratin type I, Cytoskeletal 18 (Cytokeratin 18, CK18,CYK18)) pseudogene, a gene for a novel protein, the gene for spermatogenesis associated protein PD1 (KIAA0... /FEA=mRNA_3 /DB_XREF=gi:9368423 /UG=Hs.12785 KIAA0939 protein	AL031685		
212952_at	0.042466	calreticulin	BE251303	Hs.16488	NP_004334
212953_x_at	0.009907	calreticulin	BE251303	Hs.16488	NP_004334
212958_x_at	0.003171	peptidylglycine alpha-amidating monooxygenase	AI022882	Hs.83920	NP_620177
212961_x_at	6.68E-04	Consensus includes gb:L43578.1 /DEF=Homo sapiens (clone 115392) mRNA. /FEA=mRNA /DB_XREF=gi:899067 /UG=Hs.82171 Homo sapiens (clone 115392) mRNA	L43578		NP_835225
212971_at	0.007779	cysteinyl-tRNA synthetase	AI769685	Hs.159604	NP_644802
212973_at	3.26E-04	ribose 5-phosphate isomerase A (ribose 5-phosphate epimerase)	AI692341	Hs.79886	NP_653164
212975_at	0.005852	Consensus includes gb:AB020677.2 /DEF=Homo sapiens mRNA for KIAA0870 protein, partial cds. /FEA=mRNA /GEN=KIAA0870 /PROD=KIAA0870 protein /DB_XREF=gi:6635136 /UG=Hs.18166 KIAA0870 protein	AB020677		
212982_at	3.26E-04	Huntingtin interacting protein H	AI621223	Hs.4014	NP_056151
212984_at	0.026891	activating transcription factor 2	BE786164	Hs.198166	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
212987_at	1.26E-04	Consensus includes gb:AL031178 /DEF=Human DNA sequence from clone 341E18 on chromosome 6p11.2-12.3. Contains a SerineThreonine Protein Kinase gene (presumptive isolog of a Rat gene) and a novel alternatively spliced gene. Contains a putative CpG island, ESTs and GSSs /FEA=mRNA_2 /DB_XREF=gi:3550040 /UG=Hs.11050 F-box only protein 9	AL137520		NP_258442
212990_at	1.26E-04	Consensus includes gb:AB020717.1 /DEF=Homo sapiens mRNA for KIAA0910 protein, partial cds. /FEA=mRNA /GEN=KIAA0910 /PROD=KIAA0910 protein /DB_XREF=gi:4240308 /UG=Hs.127416 synaptojanin 1	AB020717		NP_003886
212991_at	0.003171	Consensus includes gb:AL137520.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434C0118 (from clone DKFZp434C0118); partial cds. /FEA=mRNA /GEN=DKFZp434C0118 /PROD=hypothetical protein /DB_XREF=gi:6808183 /UG=Hs.11050 F-box only protein 9	AL137520		NP_258442
212993_at	0.004341	ESTs, Weakly similar to hypothetical protein FLJ12547 [Homo sapiens] [H.sapiens]	AA114166	Hs.381190	
213006_at	0.013223	CCAAT/enhancer binding protein (C/EBP), delta	AV655640	Hs.76722	NP_005186
213009_s_at	0.035171	Consensus includes gb:AK022701.1 /DEF=Homo sapiens cDNA FLJ12639 fis, clone NT2RM4001938, highly similar to Homo sapiens mRNA for KIAA0898 protein. /FEA=mRNA /DB_XREF=gi:10434250 /UG=Hs.8164 Mulibrey nanism	AK022701		NP_056109
213011_s_at	2.05E-04	triosephosphate isomerase 1	BF116254	Hs.83848	NP_000356

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213012_at	1.28E-04	Consensus includes gb:D42055.1 /DEF=Human mRNA for KIAA0093 gene, partial cds. /FEA=mRNA /GEN=KIAA0093 /DB_XREF=gi:577312 /UG=Hs.1565 neural precursor cell expressed, developmentally down-regulated 4	D42055		
213014_at	0.041795	Consensus includes gb:BG222394 /FEA=EST /DB_XREF=gi:12707915 /DB_XREF=est:naf75h09.x1 /CLONE=IMAGE:4170089 /UG=Hs.234249 mitogen-activated protein kinase 8 interacting protein 1 /FL=gb:AF074091.1 gb:NM_005456.1	NM_005456		NP_005447
213015_at	0.003539	ESTs	BF448315	Hs.171553	
213016_at	1.26E-04	ESTs	BF448315	Hs.171553	
213022_s_at	0.041055	Consensus includes gb:NM_007124.1 /DEF=Homo sapiens utrophin (homologous to dystrophin) (UTRN), mRNA. /FEA=CDS /GEN=UTRN /PROD=utrophin /DB_XREF=gi:6005937 /UG=Hs.251967 utrophin (homologous to dystrophin) /FL=gb:NM_007124.1	NM_007124		NP_009055
213027_at	0.007779	Sjogren syndrome antigen A2 (60kDa, ribonucleoprotein autoantigen SS-A/Ro)	AU146655	Hs.554	
213037_x_at	5.03E-04	Consensus includes gb:AJ132258.1 /DEF=Homo sapiens mRNA for staufen protein, partial. /FEA=mRNA /GEN=staufen /PROD=staufen protein /DB_XREF=gi:4572587 /UG=Hs.6113 staufen (Drosophila, RNA-binding protein)	AJ132258		NP_059348
213044_at	0.001116	Rho-associated, coiled-coil containing protein kinase 1	N22548	Hs.17820	NP_005397
213048_s_at	1.26E-04	SET translocation (myeloid leukemia-associated)	W26593	Hs.145279	
213049_at	0.040064	hypothetical protein DKFZp667F074	BG436400	Hs.378933	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213052_at	0.041055	protein kinase, cAMP-dependent, regulatory, type II, alpha	BF246917	Hs.365523	NP_004148
213056_at	1.26E-04	KIAA1013 protein	AU145019	Hs.96427	
213061_s_at	0.00161	Homo sapiens, Similar to N-terminal Asn amidase, clone MGC:29626 IMAGE:4872717, mRNA, complete cds	AA643304	Hs.351573	NP_775745
213073_at	0.028067	Consensus includes gb:AB002319.1 /DEF=Human mRNA for KIAA0321 gene, partial cds. /FEA=mRNA /GEN=KIAA0321 /DB_XREF=gi:2224582 /UG=Hs.8663 KIAA0321 protein	AB002319		
213084_x_at	0.00161	cadherin 1, type 1, E-cadherin (epithelial)	BF125158	Hs.356360	NP_000975
213086_s_at	0.016934	casein kinase 1, alpha 1	BF341845	Hs.283738	NP_001883
213095_x_at	5.03E-04	Consensus includes gb:AF299327.1 /DEF=Homo sapiens allograft inflammatory factor-1 splice variant G1 mRNA, complete cds. /FEA=CDS /PROD=allograft inflammatory factor-1 splice variantG1 /DB_XREF=gi:11611961 /UG=Hs.76364 allograft inflammatory factor 1 /FL=gb:AF299327.1	AF299327		NP_116573
213102_at	0.005852	ARP3 actin-related protein 3 homolog (yeast)	Z78330	Hs.5321	NP_005712
213109_at	0.009689	KIAA0551 protein	N25621	Hs.170204	
213111_at	0.02145	Consensus includes gb:AB023198.1 /DEF=Homo sapiens mRNA for KIAA0981 protein, partial cds. /FEA=mRNA /GEN=KIAA0981 /PROD=KIAA0981 protein /DB_XREF=gi:4589605 /UG=Hs.158135 KIAA0981 protein	AB023198		
213113_s_at	0.005852	likely ortholog of mouse embryonic epithelial gene 1	AI630178	Hs.274453	NP_060081

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213115_at	0.005852	Consensus includes gb:AL031177 /DEF=Human DNA sequence from clone 889N15 on chromosome Xq22.1-22.3. Contains part of the gene for a novel protein similar to X. laevis Cortical Thymocyte Marker CTX, the possibly alternatively spliced gene for 26S Proteasome subunit p28 (Ankyrin repea... /FEA=mRNA_3 /DB_XREF=gi:4071056 /UG=Hs.8763 Human DNA sequence from clone 889N15 on chromosome Xq22.1-22.3. Contains part of the gene for a novel protein similar to X. laevis Cortical Thymocyte Marker CTX, the possibly alternatively spliced gene for 26S Proteasome subunit p28 (Ankyrin repeat protei	AL031177		
213117_at	5.03E-04	KIAA1354 protein	AW138594	Hs.106283	NP_061335
213118_at	2.17E-04	Consensus includes gb:AL136821.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M1526 (from clone DKFZp434M1526). /FEA=mRNA /GEN=DKFZp434M1526 /PROD=hypothetical protein /DB_XREF=gi:12053152 /UG=Hs.153293 KIAA0701 protein	AL136821		
213119_at	0.003368	ESTs	AW058600	Hs.392501	
213120_at	0.009582	Consensus includes gb:AA405798 /FEA=EST /DB_XREF=gi:2063781 /DB_XREF=est:zu57f04.s1 /CLONE=IMAGE:742111 /UG=Hs.153293 KIAA0701 protein	AL136821		
213123_at	1.26E-04	cytochrome c oxidase subunit Vb	BE222709	Hs.1342	
213127_s_at	0.001911	mediator of RNA polymerase II transcription subunit MED8	BG230758	Hs.301756	NP_443109
213128_s_at	0.001116	ubiquitin protein ligase E3A (human papilloma virus E6-associated protein, Angelman syndrome)	AA527499	Hs.180686	NP_570854

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213133_s_at	0.001911	ESTs, Highly similar to GCHUH glycine cleavage system protein H precursor - human [H.sapiens]	BE908931	Hs.356054	
213137_s_at	0.042536	protein tyrosine phosphatase, non-receptor type 2	AI828880	Hs.82829	NP_536348
213145_at	0.041055	hypothetical protein MGC40195	BF001666	Hs.367956	
213154_s_at	2.05E-04	Consensus includes gb:AI934125 /FEA=EST /DB_XREF=gi:5672995 /DB_XREF=est:wn97c08.x1 /CLONE=IMAGE:2453774 /UG=Hs.17411 KIAA0699 protein	AB014599		NP_056065
213158_at	0.016934	Consensus includes gb:AA045174 /FEA=EST /DB_XREF=gi:1523376 /DB_XREF=est:zk66a07.s1 /CLONE=IMAGE:487764 /UG=Hs.16193 Homo sapiens mRNA; cDNA DKFZp586B211 (from clone DKFZp586B211)	AL049423		
213159_at	2.17E-04	Consensus includes gb:AB018348.1 /DEF=Homo sapiens mRNA for KIAA0805 protein, partial cds. /FEA=mRNA /GEN=KIAA0805 /PROD=KIAA0805 protein /DB_XREF=gi:3882330 /UG=Hs.55947 KIAA0805 protein	AB018348		NP_055797
213160_at	1.26E-04	Consensus includes gb:D86964.1 /DEF=Human mRNA for KIAA0209 gene, partial cds. /FEA=mRNA /GEN=KIAA0209 /DB_XREF=gi:1504001 /UG=Hs.17211 dedicator of cyto-kinesis 2	D86964		NP_004937
213165_at	0.00228	centrosome-associated protein 350	AI041204	Hs.92200	
213166_x_at	0.030771	Homo sapiens, clone IMAGE:4634703, mRNA, partial cds	BG332462	Hs.355815	
213173_at	1.49E-04	Consensus includes gb:AI815033 /FEA=EST /DB_XREF=gi:5426248 /DB_XREF=est:wk70c01.x1 /CLONE=IMAGE:2420736 /UG=Hs.55947 KIAA0805 protein	AB018348		NP_055797

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213193_x_at	0.00161	T cell receptor beta locus	AL559122	Hs.155664	
213198_at	1.26E-04	Consensus includes gb:AL117643.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M245 (from clone DKFZp434M245). /FEA=mRNA /DB_XREF=gi:5912233 /UG=Hs.5288 Homo sapiens mRNA; cDNA DKFZp434M245 (from clone DKFZp434M245)	AL117643		NP_064733
213208_at	0.007779	KIAA0240 protein	AI801951	Hs.196275	
213215_at	0.00228	Homo sapiens, clone IMAGE:4157286, mRNA, partial cds	AI910895	Hs.392619	
213216_at	0.031704	KIAA0459 protein	AL537463	Hs.374987	
213220_at	0.033381	six transmembrane epithelial antigen of prostate 2	AV706096	Hs.409158	
213222_at	0.040938	continued from dJ881E24.1.1 in Em:AL031683 match: proteins: Sw:P10687 Tr:Q9Z1B3 Sw:P10894; Human DNA sequence from clone RP4-654A7 on chromosome 20 Contains the 3' end of two variants of the PLCB1 gene for phospholipase C beta 1 (phosphoinositide-specific), complete sequence.	AL049593		
213225_at	1.26E-04	Consensus includes gb:AJ271832.1 /DEF=Homo sapiens mRNA for protein phosphatase 1B2 (PPM1B2 gene). /FEA=mRNA /GEN=PPM1B2 /PROD=protein phosphatase 1B2 53 kDa isoform /DB_XREF=gi:12666516 /UG=Hs.5687 protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform /FL=gb:NM_002706.1	AJ271832		NP_808908
213229_at	2.05E-04	Dicer1, Dcr-1 homolog (Drosophila)	BF590131	Hs.87889	NP_803187
213241_at	1.26E-04	Consensus includes gb:AF035307.1 /DEF=Homo sapiens clone 23785 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2661068 /UG=Hs.184697 Homo sapiens clone 23785 mRNA sequence	AF035307		
213243_at	0.016934	KIAA0532 protein	AI052003	Hs.21077	
213246_at	1.26E-04	DKFZP564F1123 protein	AI346504	Hs.275352	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213254_at	5.03E-04	KIAA1093 protein	N64803	Hs.117333	
213279_at	0.00161	Consensus includes gb:AL050217.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586I0523 (from clone DKFZp586I0523). /FEA=mRNA /DB_XREF=gi:4884458 /UG=Hs.308467 Homo sapiens mRNA; cDNA DKFZp586I0523 (from clone DKFZp586I0523)	AL050217		
213280_at	0.026891	Consensus includes gb:AK000478.1 /DEF=Homo sapiens cDNA FLJ20471 fis, clone KAT06974. /FEA=mRNA /DB_XREF=gi:7020593 /UG=Hs.301552 KIAA1039 protein	AK000478		
213288_at	1.41E-04	ESTs	AI761250	Hs.396081	
213292_s_at	0.001116	sorting nexin 13	AA908770	Hs.283881	NP_055947
213296_at	0.040938	peroxisome biogenesis factor 10	BF339133	Hs.247220	NP_722540
213301_x_at	0.001911	transcriptional intermediary factor 1	AL538264	Hs.183858	NP_003843
213305_s_at	0.013223	Consensus includes gb:L42375.1 /DEF=Homo sapiens protein phosphatase 2A B56-gamma1 (PP2A) mRNA, 3 end of cds. /FEA=mRNA /GEN=PP2A /PROD=protein phosphatase 2A B56-gamma1 /DB_XREF=gi:1000891 /UG=Hs.171734 protein phosphatase 2, regulatory subunit B (B56), gamma isoform	L42375		NP_848703
213309_at	0.004341	Consensus includes gb:AL117515.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L0735 (from clone DKFZp434L0735); partial cds. /FEA=mRNA /GEN=DKFZp434L0735 /PROD=hypothetical protein /DB_XREF=gi:5912029 /UG=Hs.54886 phospholipase C, epsilon 2	AL117515		NP_055999
213311_s_at	0.003539	KIAA1049 protein	BF000251	Hs.227835	NP_055787
213313_at	0.001116	rab6 GTPase activating protein (GAP and centrosome-associated)	AI922519	Hs.55099	NP_036329

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213314_at	0.027633	Consensus includes gb:NM_020425.1 /DEF=Homo sapiens hypothetical protein DKFZp586E1923 (DKFZP586E1923), mRNA. /FEA=CDS /GEN=DKFZP586E1923 /PROD=hypothetical protein DKFZp586E1923 /DB_XREF=gi:10092684 /UG=Hs.70769 hypothetical protein DKFZp586E1923 /FL=gb:NM_020425.1	NM_020425		NP_065158
213315_x_at	0.005411	Homo sapiens, clone MGC:10039 IMAGE:3889701, mRNA, complete cds	L43577	Hs.82171	NP_835225
213318_s_at	0.041055	apolipoprotein M	BG028844	Hs.274348	NP_542434
213327_s_at	0.004341	ubiquitin specific protease 12	AI820101	Hs.42400	
213341_at	0.004631	fem-1 homolog c (C.elegans)	AI862658	Hs.47367	NP_064562
213344_s_at	0.003171	H2A histone family, member X	H51429	Hs.147097	NP_002096
213351_s_at	0.008185	KIAA0779 protein	AI934469	Hs.362996	
213355_at	0.014715	Consensus includes gb:AI989567 /FEA=EST /DB_XREF=gi:5836448 /DB_XREF=est:ws34e03.x1 /CLONE=IMAGE:2499100 /UG=Hs.34578 alpha2,3-sialyltransferase	AK001922		NP_006091
213357_at	0.004341	Homo sapiens, Similar to CG14037 gene product, clone IMAGE:3640720, mRNA, partial cds	AV701318	Hs.356224	
213360_s_at	0.007779	Williams Beuren syndrome chromosome region 20C	AA514622	Hs.295112	NP_742017
213361_at	0.00161	tudor repeat associator with PCTAIRE 2	AW129593	Hs.283761	NP_055105
213365_at	0.033491	KIAA1504 protein	N64622	Hs.157426	
213366_x_at	0.041055	ATP synthase, H ⁺ transporting, mitochondrial F1 complex, gamma polypeptide 1	AV711183	Hs.155433	NP_005165
213374_x_at	0.041055	3-hydroxyisobutyryl-Coenzyme A hydrolase	AW000964	Hs.236642	NP_055177
213379_at	1.26E-04	Consensus includes gb:AF091086.1 /DEF=Homo sapiens clone 640 unknown mRNA, complete sequence. /FEA=mRNA /PROD=unknown /DB_XREF=gi:3860009 /UG=Hs.44563 hypothetical protein /FL=gb:NM_015697.1	AF091086		NP_056512

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213397_x_at	0.035171	ribonuclease, RNase A family, 4	AI761728	Hs.283749	NP_002928
213404_s_at	0.007779	601453992F1 NIH_MGC_66 Homo sapiens cDNA clone IMAGE:3857711 5', mRNA sequence.	BF033683		
213414_s_at	7.58E-04	ribosomal protein S19	BE259729	Hs.298262	NP_001013
213436_at	0.016275	G protein-coupled receptor; Human CB1 cannabinoid receptor (CNR1) gene, complete cds.	U73304		
213440_at	1.58E-04	RAB1A, member RAS oncogene family	AL530264	Hs.79507	NP_004152
213448_at	0.009716	metaxin 1	AI693193	Hs.247551	NP_002446
213453_x_at	0.041055	glyceraldehyde-3-phosphate dehydrogenase	BF689355	Hs.169476	NP_002037
213455_at	0.00161	hypothetical protein BC001096	W87466	Hs.352406	NP_612398
213463_s_at	0.041795	Consensus includes gb:AW300504 /FEA=EST /DB_XREF=gi:6710181 /DB_XREF=est:xs66c11.x1 /CLONE=IMAGE:2774612 /UG=Hs.44131 KIAA0974 protein	AB023191		
213465_s_at	5.03E-04	protein phosphatase 1, regulatory subunit 7	BF718769	Hs.36587	NP_002703
213478_at	8.84E-04	Consensus includes gb:AB028949.1 /DEF=Homo sapiens mRNA for KIAA1026 protein, partial cds. /FEA=mRNA /GEN=KIAA1026 /PROD=KIAA1026 protein /DB_XREF=gi:5689388 /UG=Hs.27742 KIAA1026 protein	AB028949		
213501_at	3.26E-04	acyl-Coenzyme A oxidase 1, palmitoyl	T62985	Hs.379991	NP_009223
213508_at	0.016282	ESTs, Weakly similar to R3HU6 ribosomal protein S6, cytosolic - human [H.sapiens]	AA142942	Hs.356665	
213510_x_at	7.58E-04	TL132 protein	AW194543	Hs.234573	NP_665808
213511_s_at	0.016934	Homo sapiens, clone MGC:17492 IMAGE:3453013, mRNA, complete cds	AI167164	Hs.347187	NP_789746
213513_x_at	0.00161	actin related protein 2/3 complex, subunit 2, 34kDa	BG034239	Hs.83583	NP_690601
213515_x_at	1.41E-04	hemoglobin, gamma A	AI133353	Hs.283108	NP_000175
213517_at	6.68E-04	poly(rC) binding protein 2	AW103422	Hs.63525	NP_114366
213521_at	0.02145	ESTs	AW575379	Hs.356456	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213524_s_at	0.007258	Consensus includes gb:NM_015714.1 /DEF=Homo sapiens putative lymphocyte G0G1 switch gene (G0S2), mRNA. /FEA=CDS /GEN=G0S2 /PROD=putative lymphocyte G0G1 switch gene /DB_XREF=gi:7657103 /UG=Hs.95910 putative lymphocyte G0G1 switch gene /FL=gb:NM_015714.1	NM_015714		NP_056529
213526_s_at	0.013223	uncharacterized hematopoietic stem/progenitor cells protein MDS033	BF215644	Hs.54960	NP_061977
213532_at	3.26E-04	a disintegrin and metalloproteinase domain 17 (tumor necrosis factor, alpha, converting enzyme)	AI797833	Hs.380136	
213535_s_at	0.013223	ubiquitin-conjugating enzyme E2I (UBC9 homolog, yeast)	AA910614	Hs.84285	NP_003336
213539_at	0.02145	Consensus includes gb:NM_000732.1 /DEF=Homo sapiens CD3D antigen, delta polypeptide (TiT3 complex) (CD3D), mRNA. /FEA=CDS /GEN=CD3D /PROD=CD3D antigen, delta polypeptide (TiT3 complex) /DB_XREF=gi:4502668 /UG=Hs.95327 CD3D antigen, delta polypeptide (TiT3 complex) /FL=gb:NM_000732.1	NM_000732		NP_000723
213541_s_at	0.004659	ESTs	AI351043	Hs.391794	NP_004440
213545_x_at	1.26E-04	sorting nexin 3	BE962615	Hs.12102	NP_690041
213549_at	1.26E-04	ESTs, Moderately similar to PRO0478 protein [Homo sapiens] [H.sapiens]	AI890972	Hs.355844	
213554_s_at	0.010205	hypothetical protein H41	BG257762	Hs.283690	NP_060018
213557_at	0.00228	ESTs	AW305119	Hs.374965	
213567_at	0.003171	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	BF431965	Hs.409065	
213571_s_at	0.00161	eukaryotic translation initiation factor 4E-like 3	BF516289	Hs.19122	NP_004837
213572_s_at	1.26E-04	serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1	AI554300	Hs.183583	NP_109591
213574_s_at	2.05E-04	karyopherin (importin) beta 1	AA861608	Hs.180446	NP_002256
213578_at	0.019761	bone morphogenetic protein receptor, type IA	AI678679	Hs.2534	NP_004320
213579_s_at	0.016934	E1A binding protein p300	AI459462	Hs.25272	NP_001420

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213582_at	0.021448	ATPase, Class VI, type 11A	BF439472	Hs.29189	
213588_x_at	0.033381	ribosomal protein L14	AA838274	Hs.738	NP_003964
213593_s_at	6.68E-04	transformer-2 alpha (htra-2 alpha)	AW978896	Hs.24937	
213603_s_at	0.007779	ras-related C3 botulinum toxin substrate 2 (rho family, small GTP binding protein Rac2)	BE138888	Hs.301175	NP_002863
213604_at	0.013223	ESTs, Highly similar to FINC_HUMAN Fibronectin precursor (FN) (Cold-insoluble globulin) (CIG) [H.sapiens]	AW451236	Hs.409286	
213614_x_at	0.016934	eukaryotic translation elongation factor 1 alpha 1	BE786672	Hs.181165	
213617_s_at	0.021312	DKFZP586M1523 protein	BF063896	Hs.22981	NP_056291
213619_at	0.004341	heterogeneous nuclear ribonucleoprotein H1 (H)	AV753392	Hs.245710	NP_005511
213620_s_at	0.011626	intercellular adhesion molecule 2	AA126728	Hs.347326	
213623_at	0.004341	Consensus includes gb:NM_007054.1 /DEF=Homo sapiens kinesin family member 3A (KIF3A), mRNA. /FEA=CDS /GEN=KIF3A /PROD=kinesin family member 3A /DB_XREF=gi:6857803 /UG=Hs.43670 kinesin family member 3A /FL=gb:AF041853.1 gb:NM_007054.1	NM_007054		NP_008985
213624_at	3.26E-04	acid sphingomyelinase-like phosphodiesterase	AA873600	Hs.391644	NP_006705
213632_at	0.01436	Consensus includes gb:M94065.1 /DEF=Human dihydroorotate dehydrogenase mRNA, 3 end. /FEA=mRNA /PROD=dihydroorotate dehydrogenase /DB_XREF=gi:555593 /UG=Hs.94925 dihydroorotate dehydrogenase	M94065		NP_001352
213637_at	0.005852	ATP-dependent RNA helicase	BE503392	Hs.380430	NP_689513
213642_at	1.26E-04	ribosomal protein L27	BE312027	Hs.111611	NP_000979
213646_x_at	7.58E-04	tubulin, alpha, ubiquitous	BE300252	Hs.334842	NP_006073
213649_at	2.05E-04	splicing factor, arginine/serine-rich 7, 35kDa	AA524053	Hs.184167	NP_006267
213666_at	6.68E-04	Consensus includes gb:AK026589.1 /DEF=Homo sapiens cDNA: FLJ22936 fis, clone KAT07936. /FEA=mRNA /DB_XREF=gi:10439473 /UG=Hs.90998 KIAA0128 protein; septin 2	AK026589		NP_665801
213671_s_at	0.005852	methionine-tRNA synthetase	AA621558	Hs.279946	NP_004981

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213672_at	0.004541	methionine-tRNA synthetase	AA621558	Hs.279946	NP_004981
213687_s_at	0.007779	ribosomal protein L35a	BE968801	Hs.287361	NP_000987
213694_at	0.016934	ESTs	AW027347	Hs.394035	
213698_at	0.013223	zinc finger protein 258	AI805560	Hs.301637	NP_660353
213702_x_at	1.26E-04	N-acylsphingosine amidohydrolase (acid ceramidase) 1	AI934569	Hs.75811	NP_808592
213708_s_at	7.58E-04	transcription factor-like 4	N40555	Hs.78185	NP_057640
213720_s_at	0.021435	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	AI831675	Hs.78202	NP_003063
213726_x_at	3.26E-04	tubulin, beta, 2	AA515698	Hs.251653	
213729_at	2.05E-04	formin binding protein 3	Z78308	Hs.107213	NP_060362
213735_s_at	2.05E-04	cytochrome c oxidase subunit Vb	AI557312	Hs.1342	NP_001853
213741_s_at	7.58E-04	karyopherin alpha 1 (importin alpha 5)	BF575685	Hs.343581	NP_002255
213786_at	0.026891	nudix (nucleoside diphosphate linked moiety X)-type motif 1	AI935415	Hs.153357	NP_006015
213794_s_at	0.041055	DKFZP564O092 protein	AI269117	Hs.9043	
213798_s_at	0.007779	adenylyl cyclase-associated protein	AA806142	Hs.104125	NP_006358
213812_s_at	0.02145	Consensus includes gb:AK024748.1 /DEF=Homo sapiens cDNA: FLJ21095 fis, clone CAS03846. /FEA=mRNA /DB_XREF=gi:10437113 /UG=Hs.108708 calciumcalmodulin-dependent protein kinase kinase 2, beta	AK024748		NP_757380
213817_at	1.26E-04	Consensus includes gb:AL049435.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586B0220 (from clone DKFZp586B0220). /FEA=mRNA /DB_XREF=gi:4500218 /UG=Hs.170056 Homo sapiens mRNA; cDNA DKFZp586B0220 (from clone DKFZp586B0220)	AL049435		
213826_s_at	0.033381	H3 histone, family 3A	AA292281	Hs.181307	NP_002098

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
213831_at	0.022752	Consensus includes gb:X00452.1 /DEF=Human mRNA for DC classII histocompatibility antigen alpha-chain. /FEA=mRNA /PROD=DC classII histocompatibility antigenalpha-chain /DB_XREF=gi:32265 /UG=Hs.198253 major histocompatibility complex, class II, DQ alpha 1	X00452		NP_002113
213836_s_at	1.26E-04	KIAA1001 protein	AW052084	Hs.279937	NP_055775
213850_s_at	6.68E-04	splicing factor, arginine/serine-rich 2, interacting protein	AI984932	Hs.51957	NP_004710
213851_at	0.001012	ESTs, Weakly similar to S65173 probable membrane protein YPL162c - yeast (Saccharomyces cerevisiae) [S.cerevisiae]	BG031677	Hs.394011	
213860_x_at	2.05E-04	casein kinase 1, alpha 1	AW268585	Hs.283738	NP_001883
213867_x_at	0.013223	actin, beta	AA809056	Hs.288061	NP_001092
213882_at	0.003529	beta-amyloid binding protein precursor	AA012917	Hs.333541	NP_114416
213883_s_at	0.003171	beta-amyloid binding protein precursor	AA012917	Hs.333541	NP_114416
213902_at	0.00228	N-acylsphingosine amidohydrolase (acid ceramidase) 1	AI379338	Hs.75811	NP_808592
213911_s_at	2.05E-04	H2A histone family, member Z	BF718636	Hs.119192	NP_002097
213918_s_at	5.03E-04	IDN3 protein	BF221673	Hs.225767	NP_597677
213919_at	0.034221	DnaJ (Hsp40) homolog, subfamily C, member 4	AW024467	Hs.172847	NP_005519
213922_at	0.003539	KIAA0847 protein	AW294686	Hs.356431	NP_775771
213923_at	5.03E-04	RAP2B, member of RAS oncogene family	AW005535	Hs.388235	NP_002877
213936_x_at	0.022795	ESTs, Moderately similar to hypothetical protein FLJ20294 [Homo sapiens] [H.sapiens]	AW276646	Hs.355462	
213940_s_at	0.003171	formin-binding protein 17	AU145053	Hs.408992	
213956_at	6.68E-04	centrosome-associated protein 350	AW299294	Hs.92200	NP_055625
213958_at	0.001618	CD6 antigen	AW134823	Hs.81226	NP_006716
213969_x_at	0.004341	ribosomal protein L29	BF683426	Hs.350068	
213971_s_at	0.022708	joined to JAZF1	AI924660	Hs.197803	NP_056170
213988_s_at	3.26E-04	spermidine/spermine N1-acetyltransferase	BE971383	Hs.396709	NP_002961
213995_at	0.02145	mitochondrial ATP synthase regulatory component factor B	AW195882	Hs.56155	NP_056499
214001_x_at	0.004341	ribosomal protein S10	AA320764	Hs.76230	NP_001005

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214022_s_at	0.041055	interferon induced transmembrane protein 1 (9-27)	AA749101	Hs.146360	
214032_at	6.68E-04	ESTs, Highly similar to ZA70_HUMAN Tyrosine-protein kinase ZAP-70 (70 kDa zeta-associated protein) (Syk-related tyrosine kinase) [H.sapiens]	AI817942	Hs.406272	
214033_at	0.008237	ATP-binding cassette, sub-family C (CFTR/MRP), member 6	AI084637	Hs.274260	NP_001162
214034_at	0.014208	type 1 tumor necrosis factor receptor shedding aminopeptidase regulator	BE551138	Hs.247043	NP_057526
214039_s_at	0.027633	putative integral membrane transporter	T15777	Hs.296398	NP_060877
214041_x_at	2.05E-04	ribosomal protein L37a	BE857772	Hs.296290	NP_000989
214055_x_at	6.68E-04	KIAA1096 protein	AW238632	Hs.69559	NP_055987
214080_x_at	0.004341	protein kinase C substrate 80K-H	AI815793	Hs.1432	NP_002734
214081_at	0.040064	Consensus includes gb:AF070526.1 /DEF=Homo sapiens clone 24787 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387881 /UG=Hs.13429 Homo sapiens clone 24787 mRNA sequence	AF070526		NP_065138
214085_x_at	0.02145	glioma pathogenesis-related protein	AI912583	Hs.154762	NP_008974
214088_s_at	0.004452	fucosyltransferase 3 (galactoside 3(4)-L-fucosyltransferase, Lewis blood group included)	AW080549	Hs.169238	NP_000140
214112_s_at	1.26E-04	Homo sapiens, clone MGC:10039 IMAGE:3889701, mRNA, complete cds	AA543076	Hs.82171	NP_835225
214119_s_at	0.033381	FK506 binding protein 1A, 12kDa	AI936769	Hs.380080	NP_463460
214121_x_at	0.007942	enigma (LIM domain protein)	AA086229	Hs.102948	NP_005442
214136_at	0.011788	nudix (nucleoside diphosphate linked moiety X)-type motif 13	W80642	Hs.16951	
214151_s_at	1.26E-04	phosphatidylinositol glycan, class B	AU144243	Hs.247118	NP_004846
214152_at	3.26E-04	phosphatidylinositol glycan, class B	AU144243	Hs.247118	NP_004846
214153_at	0.005411	homolog of yeast long chain polyunsaturated fatty acid elongation enzyme 2	BE467941	Hs.408991	
214173_x_at	0.03018	chromosome 19 open reading frame 2	AW514900	Hs.7943	NP_604431
214177_s_at	0.010205	hematopoietic PBX-interacting protein	AI935162	Hs.8068	NP_065385
214181_x_at	0.033381	leukocyte specific transcript 1	AI735692	Hs.88411	NP_009092
214198_s_at	3.26E-04	DiGeorge syndrome critical region gene 2	AU150824	Hs.2491	NP_005128

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214211_at	0.041055	ferritin, heavy polypeptide 1	AA083483	Hs.62954	NP_004174
214224_s_at	0.041055	protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin)	BE674061	Hs.11774	NP_006214
214246_x_at	0.016934	Misshapen/NIK-related kinase	AI859060	Hs.112028	NP_000071
214257_s_at	0.016934	SEC22 vesicle trafficking protein-like 1 (<i>S. cerevisiae</i>)	AA890010	Hs.50785	NP_004883
214259_s_at	0.026891	aldo-keto reductase family 7, member A2 (aflatoxin aldehyde reductase)	AW074911	Hs.6980	NP_003680
214266_s_at	0.046329	enigma (LIM domain protein)	AW206786	Hs.102948	NP_005442
214268_s_at	0.02145	myotubularin related protein 4	AL042220	Hs.141727	NP_004678
214273_x_at	0.039149	Conserved gene telomeric to alpha globin cluster	AV704353	Hs.19699	
214274_s_at	0.007779	acetyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Coenzyme A thiolase)	AI860341	Hs.166160	NP_001598
214280_x_at	0.013223	Consensus includes gb:X79536.1 /DEF=H.sapiens mRNA for hnRNPcore protein A1. /FEA=mRNA /PROD=hnRNPcore protein A1 /DB_XREF=gi:496897 /UG=Hs.249495 heterogeneous nuclear ribonucleoprotein A1	X79536		NP_112420
214281_s_at	7.58E-04	zinc finger protein 363	AA524525	Hs.48297	NP_056251
214288_s_at	0.00161	proteasome (prosome, macropain) subunit, beta type, 1	W86293	Hs.75748	NP_002784
214290_s_at	0.041055	H2A histone family, member O	AA451996	Hs.795	NP_003507
214299_at	0.021435	topoisomerase (DNA) III alpha	AI676092	Hs.91175	NP_004609
214305_s_at	6.68E-04	splicing factor 3b, subunit 1, 155kDa	AW003030	Hs.334826	NP_036565
214315_x_at	0.005852	calreticulin	AI348935	Hs.16488	NP_004334
214329_x_at	0.013223	tumor necrosis factor (ligand) superfamily, member 10	AW474434	Hs.83429	NP_003801
214352_s_at	3.26E-04	v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog	BF673699	Hs.351221	NP_203524
214366_s_at	6.68E-04	arachidonate 5-lipoxygenase	AA995910	Hs.89499	NP_000689
214403_x_at	1.72E-04	prostate epithelium-specific Ets transcription factor	AI307915	Hs.79414	NP_036523
214405_at	0.018693	ESTs	Z39557	Hs.389152	
214414_x_at	0.026891	hemoglobin, alpha 2	T50399	Hs.347939	NP_000549
214429_at	2.05E-04	Consensus includes gb:U47635.1 /DEF=Human D13S824E locus mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:1669390 /UG=Hs.79877 myotubularin related protein 6 /FL=gb:U47635.1	U47635		NP_004676

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214430_at	7.58E-04	Consensus includes gb:NM_000169.1 /DEF=Homo sapiens galactosidase, alpha (GLA), mRNA. /FEA=CDS /GEN=GLA /PROD=galactosidase, alpha /DB_XREF=gi:4504008 /UG=Hs.69089 galactosidase, alpha /FL=gb:BC002689.1 gb:NM_000169.1	NM_000169		NP_000160
214438_at	0.041055	Consensus includes gb:M60721.1 /DEF=Human homeobox gene, complete cds. /FEA=mRNA /DB_XREF=gi:183789 /UG=Hs.74870 H2.0 (Drosophila)-like homeo box 1 /FL=gb:NM_021958.1 gb:M60721.1	M60721		NP_068777
214439_x_at	0.0174	Consensus includes gb:AF043899.1 /DEF=Homo sapiens amphiphysin IIc1 mRNA, complete cds. /FEA=CDS /PROD=amphiphysin IIc1 /DB_XREF=gi:3064256 /UG=Hs.193163 bridging integrator 1 /FL=gb:AF043899.1	AF043899		NP_647601
214440_at	7.58E-04	Consensus includes gb:NM_000662.1 /DEF=Homo sapiens N-acetyltransferase 1 (arylamine N-acetyltransferase) (NAT1), mRNA. /FEA=CDS /GEN=NAT1 /PROD=N-acetyltransferase 1 /DB_XREF=gi:4505334 /UG=Hs.155956 N-acetyltransferase 1 (arylamine N-acetyltransferase) /FL=gb:NM_000662.1	NM_000662		NP_000653

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214470_at	0.041055	Consensus includes gb:NM_002258.1 /DEF=Homo sapiens killer cell lectin-like receptor subfamily B, member 1 (KLRB1), mRNA. /FEA=CDS /GEN=KLRB1 /PROD=killer cell lectin-like receptor subfamily B,member 1 /DB_XREF=gi:4504878 /UG=Hs.169824 killer cell lectin-like receptor subfamily B, member 1 /FL=gb:NM_002258.1 gb:U11276.1	NM_002258		NP_002249
214473_x_at	0.033381	Consensus includes gb:NM_005395.1 /DEF=Homo sapiens postmeiotic segregation increased 2-like 9 (PMS2L9), mRNA. /FEA=CDS /GEN=PMS2L9 /PROD=postmeiotic segregation increased 2-like 9 /DB_XREF=gi:4885552 /UG=Hs.301862 postmeiotic segregation increased 2-like 9 /FL=gb:NM_005395.1	NM_005395		NP_005386
214496_x_at	5.64E-04	Consensus includes gb:NM_012330.1 /DEF=Homo sapiens histone acetyltransferase (MORF), mRNA. /FEA=CDS /GEN=MORF /PROD=histone acetyltransferase /DB_XREF=gi:6912511 /UG=Hs.27590 histone acetyltransferase /FL=gb:AF119231.1 gb:NM_012330.1	NM_012330		NP_036462
214501_s_at	5.03E-04	Consensus includes gb:AF044286.1 /DEF=Homo sapiens histone macroH2A1.1 mRNA, complete cds. /FEA=CDS /PROD=histone macroH2A1.1 /DB_XREF=gi:3493530 /UG=Hs.75258 H2A histone family, member Y /FL=gb:AF044286.1	AF044286		NP_613258

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214511_x_at	7.58E-04	Consensus includes gb:L03419.1 /DEF=Human Fc-gamma receptor I B1 mRNA, complete cds. /FEA=CDS /PROD=Fc gamma receptor I /DB_XREF=gi:182460 /UG=Hs.77424 Fc fragment of IgG, high affinity Ia, receptor for (CD64) /FL=gb:L03419.1 gb:L03420.1	L03419		NP_000557
214512_s_at	0.00228	Consensus includes gb:NM_006713.1 /DEF=Homo sapiens activated RNA polymerase II transcription cofactor 4 (PC4), mRNA. /FEA=CDS /GEN=PC4 /PROD=activated RNA polymerase II transcriptioncofactor 4 /DB_XREF=gi:5729967 /UG=Hs.74861 activated RNA polymerase II transcription cofactor 4 /FL=gb:U12979.1 gb:NM_006713.1	NM_006713		NP_006704
214523_at	3.93E-04	Consensus includes gb:NM_001805.1 /DEF=Homo sapiens CCAATenhancer binding protein (CEBP), epsilon (CEBPE), mRNA. /FEA=CDS /GEN=CEBPE /PROD=CCAATenhancer binding protein (CEBP), epsilon /DB_XREF=gi:4502766 /UG=Hs.158323 CCAATenhancer binding protein (CEBP), epsilon /FL=gb:NM_001805.1	NM_001805		NP_001796
214531_s_at	2.05E-04	Consensus includes gb:AF065484.1 /DEF=Homo sapiens sorting nexin 1A (SNX1) mRNA, complete cds. /FEA=CDS /GEN=SNX1 /PROD=sorting nexin 1A /DB_XREF=gi:3152941 /UG=Hs.75283 sorting nexin 1 /FL=gb:AF065484.1	AF065484		NP_690039

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214548_x_at	1.26E-04	Consensus includes gb:AF064092.1 /DEF=Homo sapiens mutant guanine nucleotide-binding protein G(s), alpha subunit mRNA, complete cds. /FEA=CDS /PROD=mutant guanine nucleotide-binding protein G(s),alpha subunit /DB_XREF=gi:3142636 /UG=Hs.273385 guanine nucleotide binding protein (G protein), alpha stimulating activity polypeptide 1 /FL=gb:AF064092.1	AF064092		NP_536351
214553_s_at	0.003862	Consensus includes gb:NM_006628.1 /DEF=Homo sapiens cyclic AMP phosphoprotein, 19 kD (ARPP-19), mRNA. /FEA=CDS /GEN=ARPP-19 /PROD=cyclic AMP phosphoprotein, 19 kD /DB_XREF=gi:5729731 /UG=Hs.7351 cyclic AMP phosphoprotein, 19 kD /FL=gb:NM_006628.1	NM_006628		NP_006619
214561_at	0.027243	Consensus includes gb:NM_024317.1 /DEF=Homo sapiens immunoglobulin-like transcript 10 (ILT10), mRNA. /FEA=CDS /GEN=ILT10 /PROD=immunoglobulin-like transcript 10 /DB_XREF=gi:13399319 /UG=Hs.202680 immunoglobulin-like transcript 10 /FL=gb:NM_024317.1	NM_024317		NP_077293
214572_s_at	0.038947	Consensus includes gb:NM_005543.1 /DEF=Homo sapiens insulin-like 3 (Leydig cell) (INSL3), mRNA. /FEA=CDS /GEN=INSL3 /PROD=insulin-like 3 (Leydig cell) /DB_XREF=gi:5031802 /UG=Hs.37062 insulin-like 3 (Leydig cell) /FL=gb:NM_005543.1	NM_005543		NP_005534

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214574_x_at	0.007779	Consensus includes gb:NM_007161.1 /DEF=Homo sapiens DNA segment on chromosome 6 (unique) 49 expressed sequence, NK cell triggering receptor, p30 (D6S49E), mRNA. /FEA=CDS /GEN=D6S49E /PROD=leukocyte-specific transcript 1 /DB_XREF=gi:6005740 /UG=Hs.88411 lymphocyte antigen 117 /FL=gb:NM_007161.1	NM_007161		NP_009092
214583_at	0.012367	regulatory solute carrier protein, family 1, member 1	AI268381	Hs.239459	NP_006502
214590_s_at	2.05E-04	ubiquitin-conjugating enzyme E2D 1 (UBC4/5 homolog, yeast)	AL545760	Hs.129683	NP_003329
214594_x_at	0.005852	ATPase, Class I, type 8B, member 1	BG252666	Hs.388624	NP_005594
214606_at	0.001303	tetraspan 2	BF129969	Hs.234863	NP_005716
214615_at	0.003539	Consensus includes gb:NM_014499.1 /DEF=Homo sapiens putative purinergic receptor (P2Y10), mRNA. /FEA=CDS /GEN=P2Y10 /PROD=putative purinergic receptor /DB_XREF=gi:10092632 /UG=Hs.296433 putative purinergic receptor /FL=gb:NM_014499.1	NM_014499		NP_055314
214617_at	0.004341	perforin 1 (pore forming protein)	AI445650	Hs.2200	NP_005032
214620_x_at	3.56E-04	peptidylglycine alpha-amidating monooxygenase	BF038548	Hs.83920	NP_620177
214629_x_at	0.001116	Consensus includes gb:AF320999.1 /DEF=Homo sapiens Nogo-A protein short form mRNA, complete cds, alternatively spliced. /FEA=CDS /PROD=Nogo-A protein short form /DB_XREF=gi:11878277 /UG=Hs.65450 reticulon 4 /FL=gb:AF320999.1	AF320999		NP_722550

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214651_s_at	0.011725	Consensus includes gb:U41813.1 /DEF=Human class I homeoprotein (HOXA9) mRNA, partial cds. /FEA=mRNA /GEN=HOXA9 /PROD=HOXA9 /DB_XREF=gi:1184168 /UG=Hs.127428 homeo box A9 /FL=gb:NM_002142.1	U41813		NP_689952
214658_at	0.00228	CGI-109 protein	BG286537	Hs.278391	
214663_at	0.00228	Consensus includes gb:AB007941.1 /DEF=Homo sapiens mRNA for KIAA0472 protein, partial cds. /FEA=mRNA /GEN=KIAA0472 /PROD=KIAA0472 protein /DB_XREF=gi:3413905 /UG=Hs.6874 KIAA0472 protein	AB007941		
214665_s_at	1.26E-04	Consensus includes gb:AK000095.1 /DEF=Homo sapiens cDNA FLJ20088 fis, clone COL03869. /FEA=mRNA /DB_XREF=gi:7019960 /UG=Hs.85301 calcium binding protein P22	AK000095		
214669_x_at	0.007052	immunoglobulin kappa constant	BG485135	Hs.306357	
214684_at	5.03E-04	Consensus includes gb:X63381.1 /DEF=Homo sapiens mRNA for serum response factor-related protein, RSRFC4. /FEA=mRNA /GEN=RSRFC4 /PROD=serum response factor-related protein /DB_XREF=gi:432655 /UG=Hs.182280 MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A)	X63381		NP_005578
214687_x_at	0.033381	Consensus includes gb:AK026577.1 /DEF=Homo sapiens cDNA: FLJ22924 fis, clone KAT06977, highly similar to HSALDAR Human fibroblast mRNA for aldolase A. /FEA=mRNA /DB_XREF=gi:10439461 /UG=Hs.273415 aldolase A, fructose-bisphosphate	AK026577		NP_000025
214691_x_at	0.002704	hypothetical protein KIAA1164	AU121431	Hs.40719	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214693_x_at	0.029453	hypothetical protein DJ328E19.C1.1	BE732345	Hs.323463	NP_056198
214696_at	0.033381	Consensus includes gb:AF070569.1 /DEF=Homo sapiens clone 24659 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387938 /UG=Hs.29206 Homo sapiens clone 24659 mRNA sequence	AF070569		NP_116284
214703_s_at	7.58E-04	KIAA0935 protein	AW954107	Hs.12183	
214714_at	0.003741	Consensus includes gb:AK022360.1 /DEF=Homo sapiens cDNA FLJ12298 fis, clone MAMMA1001837, weakly similar to ZINC FINGER PROTEIN 29. /FEA=mRNA /DB_XREF=gi:10433741 /UG=Hs.284168 hypothetical protein MPMGp800M05499Q3	AK022360		NP_115540
214715_x_at	0.016934	Consensus includes gb:AK024789.1 /DEF=Homo sapiens cDNA: FLJ21136 fis, clone CAS07469. /FEA=mRNA /DB_XREF=gi:10437175 /UG=Hs.206882 Homo sapiens mRNA for FLJ00032 protein, partial cds	AK024789		NP_150630
214718_at	6.68E-04	Consensus includes gb:AK026142.1 /DEF=Homo sapiens cDNA: FLJ22489 fis, clone HRC10951. /FEA=mRNA /DB_XREF=gi:10438897 /UG=Hs.21145 hypothetical protein FLJ22489	AK026142		NP_066990
214719_at	0.010205	Consensus includes gb:AK026720.1 /DEF=Homo sapiens cDNA: FLJ23067 fis, clone LNG04993. /FEA=mRNA /DB_XREF=gi:10439638 /UG=Hs.117167 Homo sapiens cDNA: FLJ23067 fis, clone LNG04993	AK026720		

Gene Identifier	p-value	Description	Gene Accession No.	Unigen Accession No.	Protein Accession No.
214730_s_at	0.013223	Consensus includes gb:AK025457.1 /DEF=Homo sapiens cDNA: FLJ21804 fis, clone HEP00746, highly similar to HSU64791 Human Golgi membrane sialoglycoprotein MG160 (GLG1) mRNA. /FEA=mRNA /DB_XREF=gi:10437977 /UG=Hs.78979 Golgi apparatus protein 1	AK025457		NP_036333
214733_s_at	4.42E-04	Consensus includes gb:AL031427 /DEF=Human DNA sequence from clone 167A19 on chromosome 1p32.1-33. Contains three genes for novel proteins, the DIO1 gene for type I iodothyronine deiodinase (EC 3.8.1.4, TXDI1, ITDI1) and an HNRNP A3 (Heterogenous Nuclear Ribonucleoprotein A3, FBRNP) ... /FEA=mRNA_6 /DB_XREF=gi:4835258 /UG=Hs.11923 hypothetical protein	AL031427		
214737_x_at	0.005852	heterogeneous nuclear ribonucleoprotein C (C1/C2)	AV725195	Hs.182447	NP_112604
214743_at	0.003171	ESTs, Highly similar to CUT1_HUMAN CCAAT displacement protein (CDP) (Cut-like 1) [H.sapiens]	BE046521	Hs.381906	NP_001904
214748_at	0.016934	Consensus includes gb:U50529.1 /DEF=Human BRCA2 region, mRNA sequence CG016. /FEA=mRNA /DB_XREF=gi:1531600 /UG=Hs.112434 Novel human gene mapping to chromosome 13	U50529		NP_149102
214766_s_at	0.013223	Consensus includes gb:AL080144.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434N093 (from clone DKFZp434N093); partial cds. /FEA=mRNA /GEN=DKFZp434N093 /PROD=hypothetical protein /DB_XREF=gi:5262592 /UG=Hs.33363 DKFZP434N093 protein	AL080144		NP_056261

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214773_x_at	0.009689	similar to CG9578 gene product	AI983505	Hs.137576	NP_690866
214783_s_at	0.005852	annexin A11	BG177920	Hs.75510	NP_665876
214785_at	0.005852	Consensus includes gb:AB023203.1 /DEF=Homo sapiens mRNA for KIAA0986 protein, partial cds. /FEA=mRNA /GEN=KIAA0986 /PROD=KIAA0986 protein /DB_XREF=gi:4589615 /UG=Hs.53542 KIAA0986 protein	AB023203		NP_150648
214791_at	0.033381	Consensus includes gb:AK023116.1 /DEF=Homo sapiens cDNA FLJ13054 fis, clone NT2RP3001527, highly similar to Human Sp140 protein (Sp140) mRNA. /FEA=mRNA /DB_XREF=gi:10434889 /UG=Hs.158761 Homo sapiens cDNA FLJ13054 fis, clone NT2RP3001527, highly similar to Human Sp140 protein (Sp140) mRNA	AK023116		NP_612411
214798_at	0.006698	KIAA0703 gene product	AW291664	Hs.6168	
214801_at	0.004798	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	W88821	Hs.408958	
214805_at	2.05E-04	Consensus includes gb:U79273.1 /DEF=Human clone 23933 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710239 /UG=Hs.239483 Human clone 23933 mRNA sequence	U79273		
214808_at	0.001747	ESTs, Moderately similar to 2211404A B219/OB receptor [Homo sapiens] [H.sapiens]	AU147851	Hs.194397	
214812_s_at	3.26E-04	Consensus includes gb:D80006.1 /DEF=Human mRNA for KIAA0184 gene, partial cds. /FEA=mRNA /GEN=KIAA0184 /DB_XREF=gi:1136427 /UG=Hs.322903 KIAA0184 protein	D80006		
214813_at	0.001012	ESTs, Moderately similar to A48913 KRAB zinc finger protein 75 - human [H.sapiens]	W90796	Hs.389646	NP_009062
214814_at	0.007052	KIAA1966 protein	BF592058	Hs.158184	

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214816_x_at	0.021244	Consensus includes gb:BC003535.1 /DEF=Homo sapiens, clone IMAGE:3609326, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3609326) /DB_XREF=gi:13097635 /UG=Hs.192127 Homo sapiens, clone IMAGE:3609326, mRNA, partial cds	BC003535		NP_689479
214843_s_at	0.003529	Consensus includes gb:AK022864.1 /DEF=Homo sapiens cDNA FLJ12802 fis, clone NT2RP2002124, weakly similar to UBIQUITIN CARBOXYL TERMINAL HYDROLASE 4 (EC 3.1.2.15). /FEA=mRNA /DB_XREF=gi:10434503 /UG=Hs.173694 KIAA1097 protein	AK022864		NP_055832
214848_at	0.036254	Consensus includes gb:U79277.1 /DEF=Human clone 23548 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710245 /UG=Hs.71848 Human clone 23548 mRNA sequence	U79277		
214853_s_at	0.003171	SHC (Src homology 2 domain containing) transforming protein 1	AI091079	Hs.81972	NP_003020
214857_at	0.00884	Consensus includes gb:AL050035.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566H0124 (from clone DKFZp566H0124). /FEA=mRNA /DB_XREF=gi:4884276 /UG=Hs.133130 Homo sapiens mRNA; cDNA DKFZp566H0124 (from clone DKFZp566H0124)	AL050035		
214866_at	0.001911	Consensus includes gb:X74039.1 /DEF=H.sapiens mRNA for urokinase plasminogen activator receptor. /FEA=mRNA /PROD=urokinase plasminogen activator receptor /DB_XREF=gi:456192 /UG=Hs.179657 plasminogen activator, urokinase receptor	X74039		NP_002650
214875_x_at	0.016934	amyloid beta (A4) precursor-like protein 2	AW001847	Hs.279518	NP_001633

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214895_s_at	0.010205	a disintegrin and metalloproteinase domain 10	AU135154	Hs.172028	NP_001101
214902_x_at	0.007779	Consensus includes gb:AL080232.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586A061 (from clone DKFZp586A061). /FEA=mRNA /DB_XREF=gi:5262725 /UG=Hs.220696 Homo sapiens mRNA; cDNA DKFZp586A061 (from clone DKFZp586A061)	AL080232		
214906_x_at	0.004341	Consensus includes gb:AL049786.1 /DEF=Novel human gene mapping to chromosome 13. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:4902708 /UG=Hs.22174 Novel human gene mapping to chromosome 13	AL049786		NP_438169
214909_s_at	0.005339	Consensus includes gb:AK026191.1 /DEF=Homo sapiens cDNA: FLJ22538 fis, clone HRC13212, highly similar to AF070667 Homo sapiens NG,NG-dimethylarginine dimethylaminohydrolase homolog mRNA. /FEA=mRNA /DB_XREF=gi:10438961 /UG=Hs.247362 dimethylarginine dimethylaminohydrolase 2	AK026191		NP_039268
214911_s_at	6.68E-04	Consensus includes gb:S78771.1 /DEF=NAT=CpG island-associated gene human, mRNA, 1741 nt. /FEA=mRNA /DB_XREF=gi:244232 /UG=Hs.75243 bromodomain-containing 2	S78771		NP_005095
214915_at	0.005339	Consensus includes gb:AK021842.1 /DEF=Homo sapiens cDNA FLJ11780 fis, clone HEMBA1005931, weakly similar to ZINC FINGER PROTEIN 83. /FEA=mRNA /DB_XREF=gi:10433110 /UG=Hs.279935 Homo sapiens cDNA FLJ11780 fis, clone HEMBA1005931, weakly similar to ZINC FINGER PROTEIN 83	AK021842		

Gene Identifier	p-value	Description	Gene Accession No.	Unigene Accession No.	Protein Accession No.
214917_at	6.79E-04	Consensus includes gb:AK024252.1 /DEF=Homo sapiens cDNA FLJ14190 fis, clone NT2RP2006534, moderately similar to 5-AMP-ACTIVATED PROTEIN KINASE, CATALYTIC ALPHA-1 CHAIN (EC 2.7.1.-). /FEA=mRNA /DB_XREF=gi:10436581 /UG=Hs.288546 Homo sapiens cDNA FLJ14190 fis, clone NT2RP2006534, moderately similar to 5-AMP-ACTIVATED PROTEIN KINASE, CATALYTIC ALPHA-1 CHAIN (EC 2.7.1.-)	AK024252		
214919_s_at	7.58E-04	FLJ20288 protein	R39094	Hs.84045	NP_003723
214938_x_at	2.05E-04	Consensus includes gb:AF283771.2 /DEF=Homo sapiens clone TCBAP0774 mRNA sequence. /FEA=mRNA /DB_XREF=gi:10281740 /UG=Hs.274472 high-mobility group (nonhistone chromosomal) protein 1	AF283771		NP_002119
214947_at	7.10E-04	Consensus includes gb:AF052146.1 /DEF=Homo sapiens clone 24653 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360455 /UG=Hs.155085 Homo sapiens clone 24653 mRNA sequence	AF052146		
214953_s_at	2.05E-04	Consensus includes gb:X06989.1 /DEF=Human mRNA for amyloid A4(751) protein. /FEA=mRNA /DB_XREF=gi:28720 /UG=Hs.177486 amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	X06989		NP_000475
214962_s_at	0.03002	Consensus includes gb:AK026236.1 /DEF=Homo sapiens cDNA: FLJ22583 fis, clone HSI02596. /FEA=mRNA /DB_XREF=gi:10439023 /UG=Hs.22559 KIAA0197 protein	AK026236		
214964_at	0.031948	ESTs, Weakly similar to hypothetical protein FLJ20489 [Homo sapiens] [H.sapiens]	AA554430	Hs.269421	